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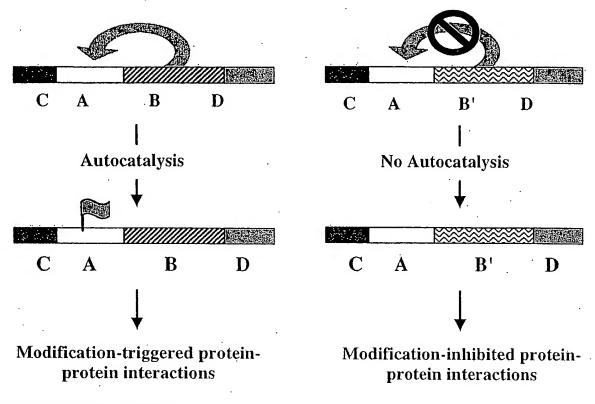
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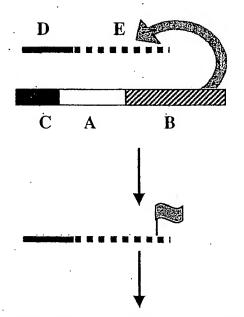
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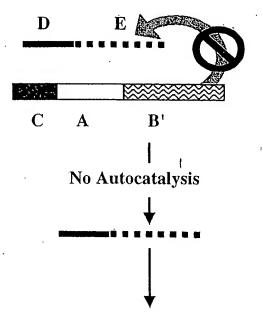
Two additional controls:



D



Modification-triggered protein-RNA interactions



Modification-inhibited protein-RNA interactions

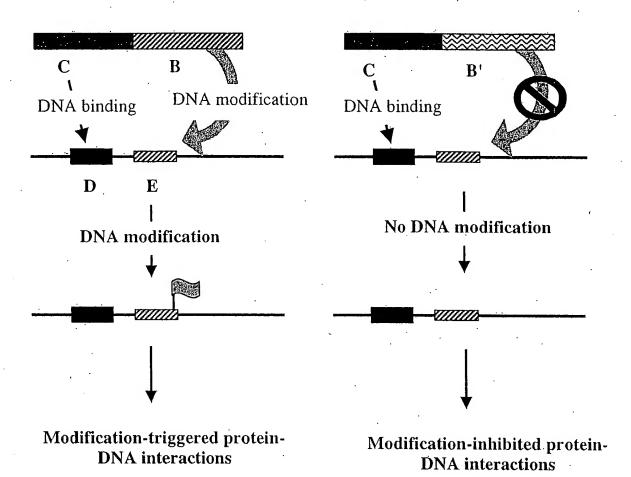
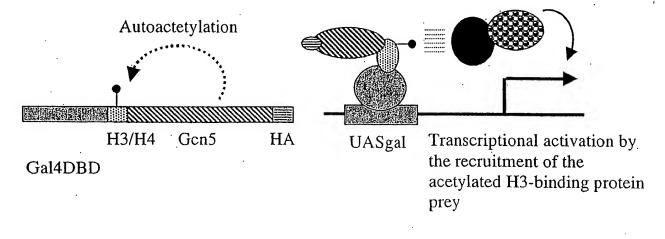
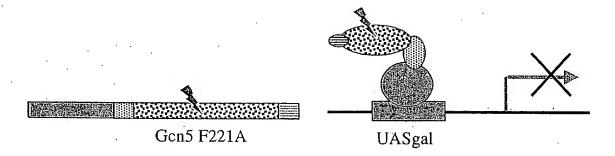


FIG: 3

Acetylation-induced interaction

Activation domain fusion prey





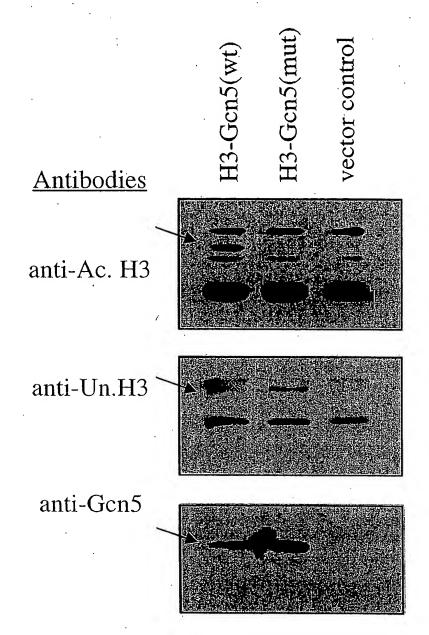
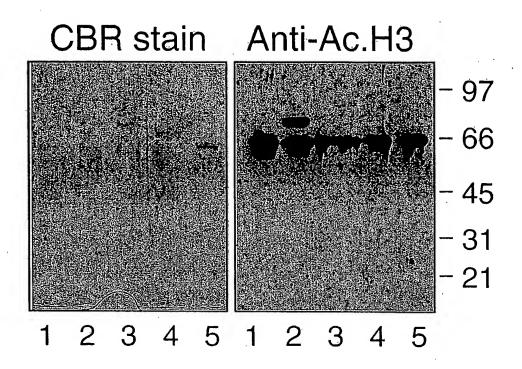


FIG. 5A



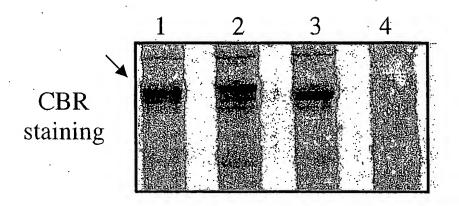
Lane 1. Vector control

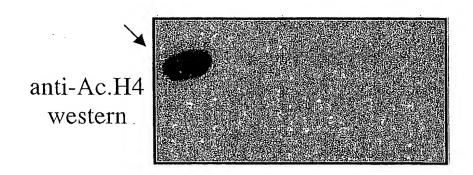
Lane 2. H3-GST-Gcn5(wt)-Ras

Lane 3. H3-GST-Gcn5(mut)-Ras

Lane 4. GST-Gcn5(wt)-Ras

Lane 5. BSA, 150 ng control

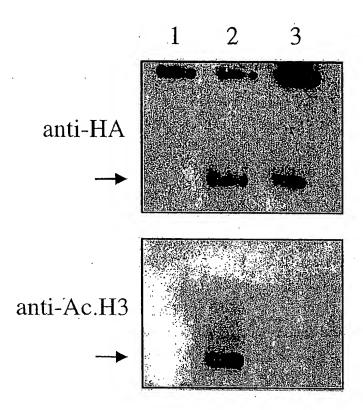




Lane 1. H4-GST-Gcn5(wt)-Ras Lane 2. H4-GST-Gcn5(mut)-Ras

Lane 3. GST-Gcn5(wt)-Ras

Lane 4. Vector control (Ras only)



Lane 1. Vector control (GDBD only) Lane 2. GDBD-H3-Gcn5(wt)-HA Lane 3. GDBD-H3-Gcn5(mut)-HA

- 1. H3-Gcn5 (wt) + vector
- 3. H4-Gcn5 (wt) + vector
- 5. H3-Gcn5 (F221A) + vector
- 7. H4-Gcn5 (F221A) + vector
- 2. H3-Gcn5 (wt) + PCAF
- 4. H4-Gcn5 (wt) + PCAF
- 6. H3-Gcn5 (F221A) + PCAF
- 8. H4-Gcn5 (F221A) + PCAF

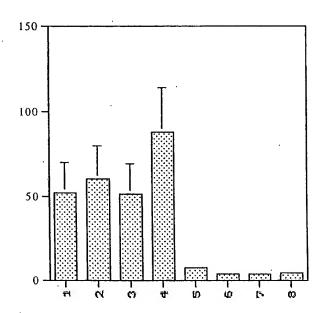


FIG. 7

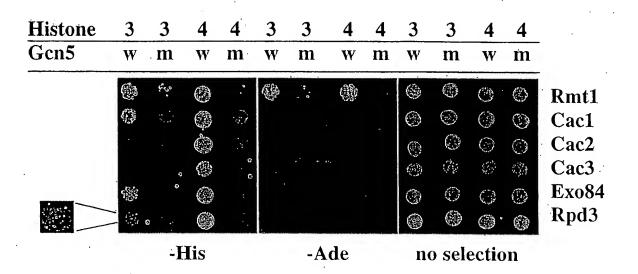


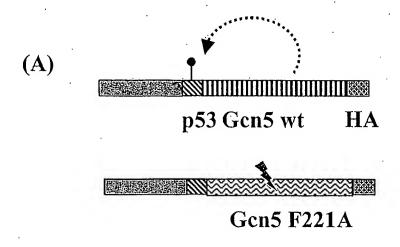
FIG. 8

Clone# 1 5 6 1 5 6

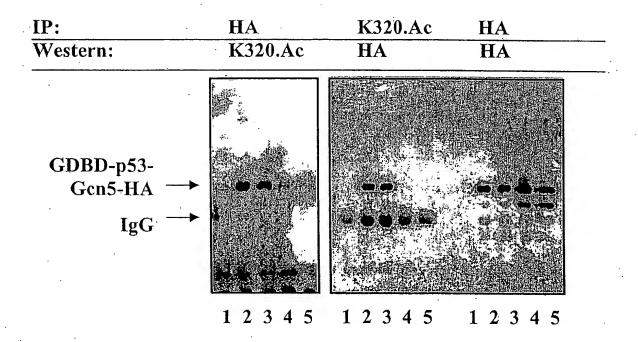
H3-Gcn5(wt)
H3-Gcn5(F221A)
H4-Gcn5(F221A)
Gcn5(F221A)
Gcn5(F221A)

-Ade no selection

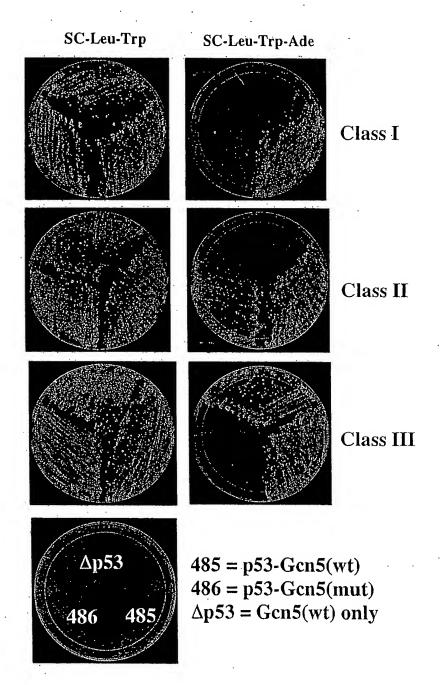
FIG. 9



(B)



- 1. Antibody only control
- 2. Wildtype Gcn5
- 3. Wildtype Gcn5
- 4. Mutant Gcn5
- 5. Mutant Gcn5



434-2059: Gal4 DBD-H3-Gcn5-HA 434-874: Gal4 DBD (amino acids 1-147) 953-1129: H3 (amino acids 1-59) 1208-1912: Gcn5 (amino acids 18-252) 1946-2035: trimeric HA: CEN/ARS ...: TRP1

Translation: Gal4-DBD-H3-Gcn5-HA

MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESRLERLEQLFLLIFPREDLDMI

LKMDSLQDIKALLTGLFVQDNVNKDAVTDRLASVETDMPLTLRQHRISATSSSEESSNKGQRQLTVSNYLFDDEDTPPNP

KKEIEFQLTTMFMARTKQTARKSTGGKAPRKQLASKAARKSAPSTGGVKKPHRYKPGTVALREIRRFQKSTEPGSPILGY

WKGRRDHPPKSDLIEGRGDPEVKRVKLENNVEEIQPEQAETNKQEGTDKENKGKFEKETERIGGSEVVTDVEKGIVKFEF

DCVEVTEKERPSVVEENEGKIEFRVVNNDNTKENMMVLTGLKNIFOKOLPKMPKEYIARLVYDRSHLSMAVIRKPLTVVG

DGVEYTFKERPSVVEENEGKIEFRVVNNDNTKENMMVLTGLKNIFQKQLPKMPKEYIARLVYDRSHLSMAVIRKPLTVVG GITYRPFDKREFAEIVFCAISSTEQVRGYGAHLMNHLKDYVRNTSNIKYFLTYADNYAIGYFKKQGFTKEITLDKSIWMG YIKDYEGGTLMQCNMAIPGGGRIFYPYDVPDYAGYPYDVPDYAGSYPYDVPDYAAQCGRSS

; DNA sequence PDG1 8285 b.p. complete sequence

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434-2059: Gal4 DBD-H3-Gcn5 (F221A)-HA 434-874: Gal4 DBD (amino acids 1-147) 953-1129: H3 (amino acids 1-59) 1208-1912: Gcn5 (amino acids 18-252) 1817-1819: F221A (1817TTT changed to GCT) 1945-2035: trimeric HA

Translation: Gal4-DBD-H3-Gcn5-HA
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gacgaaagggcctcgtgatacgcctatttttataggttaatgtcatgataataatggtttcttagacgtcaggtggcact tttcggggaaatgtgcgcggaacccctatttgtttattttctaaatacattcaaatatgtatccgctcatgagacaata accctgataaatgcttcaataatattgaaaaaggaagagtATGAGTATTCAACATTTCCGTGTCGCCCTTATTCCCTTTT TTGCGGCATTTTGCCTCCTGTTTTTGCTCACCCAGAAACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCA ${\tt CGAGTGGGTTACATCGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTTTTCGCCCCGAAGAACGTTTTCCAATGAT}$ GAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTATTATCCCGTATTGACGCCGGGCAAGAGCAACTCGGTCGCCGCATAC ACTATTCTCAGAATGACTTGGTTGAGTACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTA GTTTATTGCTGATAAATCTGGAGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGCCCT ${\tt CCCGTATCGTAGTTATCTACACGACGGGGGGTCAGGCAACTATGGATGAACGAAATAGACAGATCGCTGAGATAGGTGCC}$ taaaaggatctaggtgaagatcctttttgataatctcatgaccaaaatcccttaacgtgagttttcgttccactgagcgtccaccgctaccagcggttgtttgtttgccggatcaagagctaccaactctttttccgaaggtaactggcttcagcagagc gcagataccaaatactgtccttctagtgtagccgtagttaggccaccacttcaagaactctgtagcaccgcctacatacc 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gctttcaattcatttgggtgtgcactttattatgttacaatatggaagggaactttacacttctcctatgcacatatatt}$ aattaaagtccaatgctagtagagaggggggtaacacccctccgcgctcttttccgattttttctaaaccgtggaata $\verb|tttcggatatccttttgttgtttccgggtgtacaatatggacttcctcttttctggcaaccaaacccatacatcgggatt|\\$ cctataataccttcgttggtctccctaacatgtaggtggcggaggggagatatacaatagaacagataccagacaagaca taatgggctaaacaagactacaccaattacactgcctcattgatg

434-1963: Gal4 DBD-H4-Gcn5-HA
434-874: Gal4 DBD (amino acids 1-147)
947-1033: H4 (amino acids 1-29)
1112-1816: Gcn5 (amino acids 18-252)
1850-1939: trimeric HA
.....: CEN/ARS
.....: TRP1

Translation: Gal4 DBD-H4-Gcn5-HA
MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESRLERLEQLFLLIFPREDLDMI
LKMDSLQDIKALLTGLFVQDNVNKDAVTDRLASVETDMPLTLRQHRISATSSSEESSNKGQRQLTVSNYLFDDEDTPPNP
KKEIEFQLTTMSGRGKGGKGLGKGGAKRHRKILRDNIQGISGSPILGYWKGRRDHPPKSDLIEGRGDPEVKRVKLENNVE
EIQPEQAETNKQEGTDKENKGKFEKETERIGGSEVVTDVEKGIVKFEFDGVEYTFKERPSVVEENEGKIEFRVVNNDNTK
ENMMVLTGLKNIFQKQLPKMPKEYIARLVYDRSHLSMAVIRKPLTVVGGITYRPFDKREFAEIVFCAISSTEQVRGYGAH
LMNHLKDYVRNTSNIKYFLTYADNYAIGYFKKQGFTKEITLDKSIWMGYIKDYEGGTLMQCSMAIPGGGRIFYPYDVPDY
AGYPYDVPDYAGSYPYDVPDYAAQCGRSS

; DNA sequence PDG3 8189 b.p. complete sequence

gtggtacataacgaactaatactgtagccctagacttgatagccatcatcatatcgaagtttcactaccctttttccatt ccatatccgcaatgacaaaaaaatgatggaagacactaaaggaaaaaattaacgacaaaagacagcaccaacagatgtcgt gtttcctcgtcattgttctcgttccctttcttccttgtttcttttttctgcacaatatttcaagctataccaagcatacaa $\verb|tcaactccaagcttgaagcatcctgaaagATGAAGCTACTGTCTTCTATCGAACAAGCATGCGATATTTGCCGACT|$ TAAAAAGCTCAAGTGCTCCAAAGAAAAACCGAAGTGCGCCAAGTGTCTGAAGAACAACTGGGAGTGTCGCTACTCCCCA AAACCAAAAGGTCTCCGCTGACTAGGGCACATCTGACAGAAGTGGAATCAAGGCTAGAAAGACTGGAACAGCTATTTCTA CTGATTTTTCCTCGAGAAGACCTTGACATGATTTTGAAAATGGATTCTTTACAGGATATAAAAGCATTGTTAACAGGATT GACAGCATAGAATAAGTGCGACATCATCATCGGAAGAGAGTAGTAACAAAGGTCAAAGACAGTTGACTGTATCGAACTAT AGGTGGTAAAGGTCTAGGAAAAGGTGGTGCCAAGCGTCACAGAAAGATTCTAAGAGATAACATCCAAGGTATTTCCgggt cccctatactaggttattggaaaggtcgacgcgaccatcctccaaaatcggatctgatcgaaggtcgtggaGATCCCGAA GTTAAACGGGTAAAATTAGAAAACAACGTTGAAGAAATACAACCTGAGCAGGCTGAGACCAATAAACAAGAGGGCACCGA TAAAGAGAATAAAGGAAAGTTCGAGAAAGAAACTGAGAGAATAGGAGGATCTGAAGTGGTTACAGATGTGGAAAAAGGAA TTGTCAAATTTGAATTTGATGGTGTTGAATACACATTCAAAGAGAGACCCAGTGTCGTAGAGGAAAATGAAGGTAAAATT ${\tt GAGTTTAGGGTGGTGAATAATGATAATACTAAAGAAAACATGATGGTCCTAACTGGATTAAAAAAACATTTTTCAAAAGCA}$ ATTACCAAAAATGCCCAAAGAATACATTGCCAGGTTAGTCTATGATCGAAGTCATCTTTCCATGGCTGTCATTAGGAAGC CATTGACTGTCGTAGGTGGCATAACATATCGACCTTTCGATAAGAGAGAATTCGCAGAAATTGTTTTCTGTGCCATCAGT TCGACGGAACAGGTACGCGGTTATGGTGCGCATCTAATGAATCACTTAAAAGACTATGTTAGAAAATACCTCGAACATAAA ATATTTTTTGACATATGCAGATAATTACGCTATTGGATACTTTAAAAAGCAAGGCTTTACTAAAAGAAATCACGTTGGATA cqcatcttttacccatacgatgttcctgactatgcgggctatccctatgacgtcccggactatgcaggatcctatccata tacatcgttttgccttcttttatgtaactatactcctctaagtttcaatcttggccatgtaacctctgatctatagaattttttaaatgactagaattaatgcccatcttttttttggacctaaattcttcatgaaaatatattacgagggcttattcag a agettt ggacttettegee ag aggttt ggte aag tetee aate aaggttg teggettg tetaeettgee ag aa atttaegaaaagatggaaaagggtcaaatcgttggtagatacgttgttgacacttctaaataagcgaatttcttatgatttatgat ttttattattaaataagttataaaaaaaataagtgtatacaaattttaaagtgactcttaggttttaaaacgaaaattct 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TTTATATGCTTTTACAAGACTTGAAATTTTCCTTGCAATAACCGGGTCAATTGTTCTCTTTTCTATTGGGCACACATATAA TACCCAGCAAGTCAGCATCGGAATCTAGTGCACATTCTGCGGCCTCTGTGCTCTGCAAGCCGCAAACTTTCACCAATGGA CCAGAACTACCTGTGAAATTAATAACAGACATactccaagctgcctttgtgtgcttaatcacgtatactcacgtgctcaa gatacgcctatttttataggttaatgtcatgataataatggtttcttagacgtcaggtggcacttttcgggggaaatgtgc gcggaacccctatttgtttatttttctaaatacattcaaatatgtatccgctcatgagacaataaccctgataaatgctt caataatattgaaaaaggaagagtATGAGTATTCAACATTTCCGTGTCGCCCTTATTCCCTTTTTTGCGGCATTTTGCCT TCCTGTTTTTGCTCACCCAGAAACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTACATCG AACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTTTTCGCCCCGAAGAACGTTTTCCAATGATGAGCACTTTTAAAGTT CTGCTATGTGGCGCGGTATTATCCCGTATTGACGCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGA CTTGGTTGAGTACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATGCAGTGCCATAA CCATGAGTGATAACACTGCGGCCAACTTACTTCTGACAACGATCGGAGGACCGAAGGAGCTAACCGCTTTTTTTGCACAAC TCTGGAGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGCCCTCCCGTATCGTAGTTAT CTACACGACGGGGAGTCAGGCAACTATGGATGAACGAAATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATT aagatcctttttgataatctcatgaccaaaatcccttaacgtgagttttcgttccactgagcgtcagaccccgtagaaaa ${\tt tggtttgtttgccggatcaagagctaccaactctttttccgaaggtaactggcttcagcagagcgcagataccaaatact}$ $\tt gtccttctagtgtagccgtagttaggccaccacttcaagaactctgtagcaccgcctacatacctcgctctgctaatcct$ gttaccagtggctgctgccagtggcgataagtcgtgtcttaccgggttggactcaagacgatagttaccggataaggcgc

agcggtcgggctgaacggggggttcgtgcacacagcccagcttggagcgaacgacctacaccgaactgagatacctacag cgtgagctatgagaaagcgccacgcttcccgaagggagaaaggcggacaggtatccggtaagcggcagggtcggaacagg agagcgcacgagggagcttccagggggaaacgcctggtatctttatagtcctgtcgggtttcgccacctctgacttgagc $\tt gtcgatttttgtgatgctcgtcagggggggggggcctatggaaaaacgccagcaacgcggcctttttacggttcctggcc$ $\verb|tttgctggccttttgctcacatgttctttcctgcgttatcccctgattctgtggataaccgtattaccgcctttgagtg|$ agctgataccgctcgccgcagccgaacgaccgagcgcagcgagtcagtgagcgaggaagcggaagagcgcccaatacgca aaccgcctctccccgcgcgttggccgattcattaatgcaggatccgggatcgaagaaatgatggtaaatgaaataggaaa tttgcggcgccgaaaaacgagtttacgcaattgcacaatcatgctgactctgtggcggacccgcgctcttgccggcccg gcgataacgctgggcgtgaggctgtgcccggcggagttttttgcgcctgcattttccaaggtttaccctgcgctaagggg cgagattggagaagcaataagaatgccggttggggttgcgatgatgacgaccacgacaactggtgtcattatttaagttg ccgaaagaacctgagtgcatttgcaacatgagtatactagaagaatgagccaagacttgcgagacgcgagtttgccggtg gtgcgaacaatagagcgaccatgaccttgaaggtgagacgcgcataaccgctagagtactttgaagaggaaacagcaata gggttgctaccagtataaatagacaggtacatacaacactggaaatggttgtctgtttgagtacgctttcaattcatttg tagtagagagggggtaacacccctccgcgctcttttccgatttttctaaaccgtggaatattccggatatccttttgttgtttccgggtgtacaatatggacttcctcttttctggcaaccaaacccatacatcgggattcctataataccttcgt ${\tt tggtctccctaacatgtaggtggcggagggagatatacaatagaacagataccagacaagacataatgggctaaacaag}$ actacaccaattacactgcctcattgatg //

434-1963: Gal4 DBD-H4-Gcn5-HA
434-874: Gal4 DBD (amino acids 1-147)
947-1033: H4 (amino acids 1-29)
1112-1816: Gcn5 (amino acids 18-252)
1721-1723: F221A point mutation
1850-1939: trimeric HA
.....: CEN/ARS
....: TRP1

Translation: Gal4 DBD-H4-Gcn5-HA
MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESRLERLEQLFLLIFPREDLDMI
LKMDSLQDIKALLTGLFVQDNVNKDAVTDRLASVETDMPLTLRQHRISATSSSEESSNKGQRQLTVSNYLFDDEDTPPNP
KKEIEFQLTTMSGRGKGGKGLGKGGAKRHRKILRDNIQGISGSPILGYWKGRRDHPPKSDLIEGRGDPEVKRVKLENNVE
EIQPEQAETNKQEGTDKENKGKFEKETERIGGSEVVTDVEKGIVKFEFDGVEYTFKERPSVVEENEGKIEFRVVNNDNTK
ENMMVLTGLKNIFQKQLPKMPKEYIARLVYDRSHLSMAVIRKPLTVVGGITYRPFDKREFAEIVFCAISSTEQVRGYGAH
LMNHLKDYVRNTSNIKYFLTYADNYAIGYAKKQGFTKEITLDKSIWMGYIKDYEGGTLMQCSMAIPGGGRIFYPYDVPDY
AGYPYDVPDYAGSYPYDVPDYAAQCGRSS

; DNA sequence PDG3 8189 b.p. complete sequence

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TAAAGAGAATAAAGGAAAGTTCGAGAAAGAAACTGAGAGAATAGGAGGATCTGAAGTGGTTACAGATGTGGAAAAAGGAA TTGTCAAATTTGAATTTGATGGTGTTGAATACACATTCAAAGAGAGACCCAGTGTCGTAGAGGAAAATGAAGGTAAAATT GAGTTTAGGGTGGTGAATAATGATAATACTAAAGAAAACATGATGGTCCTAACTGGATTAAAAAAACATTTTTCAAAAGCA ATTACCAAAAATGCCCAAAGAATACATTGCCAGGTTAGTCTATGATCGAAGTCATCTTTCCATGGCTGTCATTAGGAAGC CATTGACTGTCGTAGGTGGCATAACATATCGACCTTTCGATAAGAGAGAATTCGCAGAAATTGTTTTCTGTGCCATCAGT TCGACGGAACAGGTACGCGGTTATGGTGCGCATCTAATGAATCACTTAAAAGACTATGTTAGAAATACCTCGAACATAAA ATATTTTTTGACATATGCAGATAATTACGCTATTGGATACTTTAAAAAGCAAGGCTTTACTAAAAGAAATCACGTTGGATA cgcatcttttacccatacgatgttcctgactatgcgggctatccctatgacgtcccggactatgcaggatcctatccata ${\tt tacatcgttttgccttcttttatgtaactatactcctctaagtttcaatcttggccatgtaacctctgatctatagaatt}$ $\verb|ttttaaatgactagaattaatgcccatcttttttttggacctaaattcttcatgaaaatatattacgagggcttattcag|$ ${\tt aagctttggacttcttcgccagaggtttggtcaagtctccaatcaaggttgtcggcttgtctaccttgccagaaatttac}$ gaaaagatggaaaagggtcaaatcgttggtagatacgttgttgacacttctaaataagcgaatttcttatgatttatgat 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CACCAATAACGCCATTTAATCTAAGCGCATCACCAACATTTTCTGGCGTCAGTCCACCAGCTAACATAAAATGTAAGCTC TCGGGGCTCTCTTGCCTTCCAACCCAGTCAGAAATCGAGTTCCAATCCAAAAGTTCACCTGTCCCACCTGCTTCTGAATC AAACAAGGGAATAAACGAATGAGGTTTCTGTGAAGCTGCACTGAGTATGTTGCAGTCTTTTGGAAATACGAGTCTTT TAATAACTGGCAAACCGAGGAACTCTTGGTATTCTTGCCACGACTCATCTCCATGCAGTTGGACGATATCAATGCCGTAA TTTATATGCTTTTACAAGACTTGAAATTTTCCTTGCAATAACCGGGTCAATTGTTCTCTTTTCTATTGGGCACACATATAA TACCCAGCAAGTCAGCATCGGAATCTAGTGCACATTCTGCGGCCTCTGTGCTGCAAGCCGCAAACTTTCACCAATGGA CCAGAACTACCTGTGAAATTAATAACAGACATactccaagctgcctttgtgtgcttaatcacgtatactcacgtgctcaa gatacgcctatttttataggttaatgtcatgataataatggtttcttagacgtcaggtggcacttttcgggggaaatgtgc gcggaacccctatttgtttattttctaaatacattcaaatatgtatccgctcatgagacaataaccctgataaatgctt caataatattgaaaaaggaagagtATGAGTATTCAACATTTCCGTGTCGCCCTTATTCCCTTTTTTGCGGCATTTTGCCT TCCTGTTTTTGCTCACCCAGAAACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTACATCG ${\tt AACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTTTTCGCCCCGAAGAACGTTTTCCAATGATGAGCACTTTTAAAGTT}$ ${\tt CTGCTATGTGGCGCGGTATTATCCCGTATTGACGCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGA}$ CTTGGTTGAGTACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATGCAGTGCTGCCATAA CCATGAGTGATAACACTGCGGCCAACTTACTTCTGACAACGATCGGAGGACCGAAGGAGCTAACCGCTTTTTTGCACAAC ${\tt TCTGGAGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGCCCTCCCGTATCGTAGTTAT}$ CTACACGACGGGGAGTCAGGCAACTATGGATGAACGAAATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATT aagatootttttgataatotoatgaccaaaatooottaacgtgagttttogttocactgagogtcagaccccgtagaaaa ${\tt tggtttgtttgccggatcaagagctaccaactctttttccgaaggtaactggcttcagcagagcgcagataccaaatact}$

gtccttctagtgtagccgtagttaggccaccacttcaagaactctgtagcaccgcctacatacctcgctctgctaatcct gttaccagtggctgctgccagtggcgataagtcgtgtcttaccgggttggactcaagacgatagttaccggataaggcgc agcggtcgggctgaacggggggttcgtgcacacagcccagcttggagcgaacgacctacaccgaactgagatacctacag cgtgagctatgagaaagcgccacgcttcccgaagggagaaaggcggacaggtatccggtaagcggcagggtcggaacagg agagcgcacgagggagcttccagggggaaacgcctggtatctttatagtcctgtcgggtttcgccacctctgacttgagc $\tt gtcgatttttgtgatgctcgtcagggggggggggcctatggaaaaacgccagcaacgcggcctttttacggttcctggcc$ ttttgctggccttttgctcacatgttctttcctgcgttatcccctgattctgtggataaccgtattaccgcctttgagtg agctgataccgctcgccgcagccgaacgaccgagcgcagcgagtcagtgagcgaggaagcggaagagcgcccaatacgca aaccgcctctccccgcgcgttggccgattcattaatgcaggatccgggatcgaagaaatgatggtaaatgaaataggaaa tttgcggcgccgaaaaacgagtttacgcaattgcacaatcatgctgactctgtggcggacccgcgctcttgccggcccg gcgataacgctgggcgtgaggctgtgcccggcggagttttttgcgcctgcattttccaaggtttaccctgcgctaagggg cgagattggagaagcaataagaatgccggttggggttgcgatgatgacgaccacgacaactggtgtcattatttaagttg ccgaaagaacctgagtgcatttgcaacatgagtatactagaagaatgagccaagacttgcgagacgcgagtttgccggtg gtgcgaacaatagagcgaccatgaccttgaaggtgagacgcgcataaccgctagagtactttgaagaggaaacagcaata gggttgctaccagtataaatagacaggtacatacaacactggaaatggttgtctgtttgagtacgctttcaattcatttg tagtagagagagggggtaacacccctccgcgctcttttccgatttttctaaaccgtggaatatttcggatatccttttgttgtttccgggtgtacaatatggacttcctcttttctggcaaccaaacccatacatcgggattcctataataccttcgt tggtctccctaacatgtaggtggcggagggagatatacaatagaacagataccagacaagacataatgggctaaacaag actacaccaattacactgcctcattgatg //

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5988-7613: Gal4 DBD-H3-Gcn5-HA
5988-6428: Gal4 DBD (amino acids 1-147)
6507-6683: H3 (amino acids 1-59)
6762-7466: Gcn5 (amino acids 18-252)
7500-7589: trimeric HA
.....: 2µ
.....: LEU2
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Translation: Gal4-DBD-H3-Gcn5-HA
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KKEIEFQLTTMFMARTKQTARKSTGGKAPRKQLASKAARKSAPSTGGVKKPHRYKPGTVALREIRRFQKSTEPGSPILGY
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DGVEYTFKERPSVVEENEGKIEFRVVNNDNTKENMMVLTGLKNIFQKQLPKMPKEYIARLVYDRSHLSMAVIRKPLTVVG
GITYRPFDKREFAEIVFCAISSTEQVRGYGAHLMNHLKDYVRNTSNIKYFLTYADNYAIGYFKKQGFTKEITLDKSIWMG
YIKDYEGGTLMQCNMAIPGGGRIFYPYDVPDYAGYPYDVPDYAGSYPYDVPDYAAQCGRSS

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; ### from DNA Strider Saturday, March 22, 2003 1:27:34 PM; DNA sequence pdg5 7891 b.p. complete sequence
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ATACATTCTAATGTCTGCCCCTATGTCTGCCCCTAAGAAGATCGTCGTTTTTGCCAGGTGACCACGTTGGTCAAGATCACA TGGTGCTGCTATCGATGCTACAGGTGTCCCACTTCCAGATGAGGCGCTGGAAGCCTCCAAGAAGGTTGATGCCGTTTTGT TAGGTGCTGTGGGTGGTCCTAAATGGGGTACAGGTAGTGTTAGACCTGAACAAGGTTTACTAAAAATCCGTAAAGAACTT CAATTGTACGCCAACTTAAGACCATGTAACTTTGCATCCGACTCTCTTTTAGACTTATCTCCAATCAAGCCACAATTTGC TAAAGGTACTGACTTCGTTGTTGTCAGAGAATTAGTGGGAGGTATTTACTTTGGTAAGAGAAAGGAAGACGATGGTGATG GTGTCGCTTGGGATAGTGAACAATACACCGTTCCAGAAGTGCAAAGAATCACAAGAATGGCCGCTTTCATGGCCCTACAA CATGAGCCACCATTGCCTATTTGGTCCTTGGATAAAGCTAATGTTTTGGCCTCTTCAAGATTATGGAGAAAAACTGTGGA ACCCAACCCACCTAAATGGTATTATAATCACCAGCAACATGTTTGGTGATATCATCTCCGATGAAGCCTCCGTTATCCCA GGTTCCTTGGGTTTGTTGCCATCTGCGTCCTTGGCCTCTTTGCCAGACAAGAACACCGCATTTGGTTTGTACGAACCATG CCACGGTTCTGCTCCAGATTTGCCAAAGAATAAGGTTGACCCTATCGCCACTATCTTGTCTGCTGCAATGATGTTGAAAT TGTCATTGAACTTGCCTGAAGAAGGTAAGGCCATTGAAGATGCAGTTAAAAAAGGTTTTGGATGCAGGTATCAGAACTGGT GATTTAGGTGGTTCCAACAGTACCACCGAAGTCGGTGATGCTGTCGCCGAAGAAGTTAAGAAAATCCTTGCTTAAAAAAGA TTCTCTTTTTTTATGATATTTGTACATAAACTTTATAAATGAAATTCATAATAGAAACGACACGAAATTACAAAATGGAA CCCGCATGGAATGGGATAATATCACAGGAGGTACTAGACTACCTTCATCCTACATAAATAGACGCATATAAGTACGCAT CAGTGAGCTGTATGTGCGCAGCTCGCGTTGCATTTTCGGAAGCGCTCGTTTTCGGAAACGCTTTGAAGTTCCTATTCCGA AGTTCCTATTCTCTAGCTAGAAAGTATAGGAACTTCAGAGCGCTTTTGAAAACCAAAAGCGCTCTGAAGACGCACTTTCA AAAAACCAAAAACGCACCGGACTGTAACGAGCTACTAAAATATTGCGAATACCGCTTCCACAAACATTGCTCAAAAGTAT CTCTTTGCTATATATCTCTGTGCTATATCCCTATATAACCTACCCATCCACCTTTCGCTCCTTGAACTTGCATCTAAACT CGACCTCTACATTTTTTATGTTTATCTCTAGTATTACTCTTTAGACAAAAAATTGTAGTAAGAACTATTCATAGAGTGA ATCGAAAACAATACGAAAATGTAAACATTTCCTATACGTAGTATATAGAGACAAAATAGAAGAAAACCGTTCATAATTTTC TGACCAATGAAGAATCATCAACGCTATCACTTTCTGTTCACAAAGTATGCGCAATCCACATCGGTATAGAATATAATCGG GGATGCCTTTATCTTGAAAAAATGCACCCGCAGCTTCGCTAGTAATCAGTAAACGCGGGAAGTGGAGTCAGGCTTTTTTT ATGGAAGAGAAAATAGACACCAAAGTAGCCTTCTTCTAACCTTAACGGACCTACAGTGCAAAAAGTTATCAAGAGACTGC ATTATAGAGCGCACAAAGGAGAAAAAAGTAATCTAAGATGCTTTGTTAGAAAAATAGCGCTCTCGGGATGCATTTTTGT AGAACAAAAAAGAAGTATAGATTCTTTGTTGGTAAAATAGCGCTCTCGCGTTGCATTTCTGTTCTGTAAAAATGCAGCTC AGATTCTTTGTTTGAAAAATTAGCGCTCTCGCGTTGCATTTTTGTTTTACAAAAATGAAGCACAGATTCTTCGTTGGTAA GCATTTTTGTTCTACAAAATGAAGCACAGATGCTTCGTTAACAAAGATATGCTATTGAAGTGCAAGATGGAAACGCAGAA TCTTCCGTAAAGCGCTAGACTATATTATTATACAGGTTCAAATATACTATCTGTTTCAGGGAAAACTCCCAGGTTCGG ATGTTCAAAATTCAATGATGGGTAACAAGTACGATCCGATATATGCGGTGTGAAATACCGCACAGATGCGTAAGGAGAAA ATACCGCATCAGGCGCCATTCGCCATTCAGGCTGCGCAACTGTTGGGAAGGGCGATCGGTGCGGGCCTCTTCGCTATTAC ACGACGGCCAGTGAATTcatgtaggtggcggaggggagatatacaatagaacagataccagacaagacataatgggctaa acaagactacaccaattacactgcctcattgatggtggtacataacgaactaatactgtagccctagacttgatagccat catcatatcgaagtttcactaccctttttccatttgccatctattgaagtaataataggcgcatgcaacttcttttcttt ttttttcttttctctctcccccgttgttgtctcaccatatccgcaatgacaaaaaaatgatggaagacactaaaggaaaa aattaacgacaaagacagcaccaacagatgtcgttgttccagagctgatgaggggtatctcgaagcacacgaaacttttt ccttccttcattgacctgcaattattaatcttttgtttcctcgtcattgttctcgttccctttcttccttgtttctttt $\verb|CTTCTATCGAACAAGCATGCGATATTTGCCGACTTAAAAAGCTCAAGTGCTCCAAAGAAAAACCGAAGTGCCCCAAGTGT|\\$ CTGAAGAACAACTGGGAGTGTCGCTACTCTCCCAAAACCAAAAGGTCTCCGCTGACTAGGGCACATCTGACAGAAGTGGA ATCAAGGCTAGAAAGACTGGAACAGCTATTTCTACTGATTTTTCCTCGAGAAGACCTTGACATGATTTTGAAAATGGATT GCTTCAGTGGAGACTGATATGCCTCTAACATTGAGACAGCATAGAATAAGTGCGACATCATCATCGGAAGAGAGTAGTAA AATTCCAGCTGACCACCATGTTTatggccagaacaaagcaaacagcaagaaagtccactggtggtaaggccccaagaaag caattagcttctaaggctgccagaaaatccgccccatctaccggtggtgttaagaagcctcacagatataagccaggtac ${\tt tgttgctttgagagaaatcagaagattccaaaaatctactgaaCCCgggtcccctatactaggttattggaaaggtcgac}$ gcgaccatcctccaaaatcggatctgatcgaaggtcgtggaGATCCCGAAGTTAAACGGGTAAAATTAGAAAACAACGTT GAAGAATACAACCTGAGCAGGCTGAGACCAATAAACAAGAGGGCACCGATAAAGAGAATAAAGGAAAGTTCGAGAAAGA AACTGAGAGAATAGGAGGATCTGAAGTGGTTACAGATGTGGAAAAAGGAATTGTCAAATTTGAATTTGATGGTGTTGAAT AAAGAAAACATGATGGTCCTAACTGGATTAAAAAAACATTTTTCAAAAGCAATTACCAAAAAATGCCCAAAGAATACATTGC CAGGTTAGTCTATGATCGAAGTCATCTTTCCATGGCTGTCATTAGGAAGCCATTGACTGTCGTAGGTGGCATAACATATC GACCTTTCGATAAGAGAAATTCGCAGAAATTGTTTTCTGTGCCATCAGTTCGACGGAACAGGTACGCGGTTATGGTGCG CATCTAATGAATCACTTAAAAGACTATGTTAGAAATACCTCGAACATAAAATATTTTTTGACATATGCAGATAATTACGC ATGAAGGTGGTACGCTGATGCAATGTAACATGGCAATTCCCGGtggcggccgcatcttttacccatacgatgttcctgac ${\tt tatgcgggctatccctatgacgtcccggactatgcaggatcctatccatatgacgttccagattacgctgctcagtgcgg}$ $\verb|ccgctctagctagaactagtggatcccccGATACCGTCGACCTGCAGGCATGCAAGCTTGGCGTAATCATGGTCATAGCT|\\$ GTTTCCTGTGTGAAATTGTTATCCGCTCACAATTCCACACAACATACGAGCCGGAAGCATAAAGTGTAAAGCCTGGGGTG CCTAATGAGTGAGCTAACTCACATTAATTGCGTTGCGCTCACTGCCCGCTTTCCAGTCGGGAAACCTGTCGTGCCAGCTG CATTAATGAATCGGCCAACGCGCGGGGAGAGGCGGTTTGCGTATTGGGCGC

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5988-7613: Gal4 DBD-H3-Gcn5(F221A)-HA
5988-6428: Gal4 DBD (amino acids 1-147)
6507-6683: H3 (amino acids 1-59)
6762-7466: Gcn5 (amino acids 18-252)
7371-7373: F221A
7500-7589: trimeric HA
...... 2μ
...... LEU2
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Translation: Gal4-DBD-H3-Gcn5-HA

MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESRLERLEQLFLLIFPREDLDMI LKMDSLQDIKALLTGLFVQDNVNKDAVTDRLASVETDMPLTLRQHRISATSSSEESSNKGQRQLTVSNYLFDDEDTPPNP KKEIEFQLTTMFMARTKQTARKSTGGKAPRKQLASKAARKSAPSTGGVKKPHRYKPGTVALREIRRFQKSTEPGSPILGY WKGRRDHPPKSDLIEGRGDPEVKRVKLENNVEEIQPEQAETNKQEGTDKENKGKFEKETERIGGSEVVTDVEKGIVKFEF DGVEYTFKERPSVVEENEGKIEFRVVNNDNTKENMMVLTGLKNIFQKQLPKMPKEYIARLVYDRSHLSMAVIRKPLTVVG GITYRPFDKREFAEIVFCAISSTEQVRGYGAHLMNHLKDYVRNTSNIKYFLTYADNYAIGYAKKQGFTKEITLDKSIWMG YIKDYEGGTLMQCNMAIPGGGRIFYPYDVPDYAGYPYDVPDYAGSYPYDVPDYAAQCGRSS

; DNA sequence pdg6 7891 b.p. complete sequence

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GGAGGGCTTACCATCTGGCCCCAGTGCTGCAATGATACCGCGAGACCCACGCTCACCGGCTCCAGATTTATCAGCAATAA GAAGCTAGAGTAAGTAGTTCGCCAGTTAATAGTTTGCGCAACGTTGTTGCCATTGCTACAGGCATCGTGGTGTCACGCTC GTCGTTTGGTATGGCTTCATTCAGCTCCGGTTCCCAACGATCAAGGCGAGTTACATGATCCCCCATGTTGTGCAAAAAAAG CATAATTCTCTTACTGTCATGCCATCCGTAAGATGCTTTTCTGTGACTGGTGAGTACTCAACCAAGTCATTCTGAGAATA GTGTATGCGGCGACCGAGTTGCTCTTGCCCGGCGTCAATACGGGATAATACCGCGCCACATAGCAGAACTTTAAAAGTGC TCATCATTGGAAAACGTTCTTCGGGGCGAAAACTCTCAAGGATCTTACCGCTGTTGAGATCCAGTTCGATGTAACCCACT CGTGCACCCAACTGATCTTCAGCATCTTTTACTTTCACCAGCGTTTCTGGGTGAGCAAAAACAGGAAGGCAAAAATGCCGC AAAAAGGGAATAAGGGCGACACGGAAATGTTGAATACTCATACTCTTTCCTTTTTCAATATTATTGAAGCATTTATCAGG GTTATTGTCTCATGAGCGGATACATATTTGAATGTATTTAGAAAAATAAACAAATAGGGGTTCCGCGCACATTTCCCCGA AAAGTGCCACCTGACGTCTAAGAAACCATTATTATCATGACATTAACCTATAAAAATAGGCGTATCACGAGGCCCTTTCG TCTTCAAGAATTAACTGTGGGAATACTCAGGTATCGTAAGATGCAAGAGTTCGAATCTCTTAGCAACCATTATTTTTTTC ${\tt CGCCGCGCATATACCTTTTTCAACTGAAAAATTGGGAGAAAAAGGAAAGGTGAGAGGCCGGAACCGGCTTTTCATATAGA}$ ATAGAGAAGCGTTCATGACTAAATGCTTGCATCACAATACTTGAAGTTGACAATATTATTTAAGGACCTATTGTTTTTC ATACATTCTAATGTCTGCCCCTATGTCTGCCCCTAAGAAGATCGTCGTTTTGCCAGGTGACCACGTTGGTCAAGATCACA TGGTGCTGCTATCGATGCTACAGGTGTCCCACTTCCAGATGAGGCGCTGGAAGCCTCCAAGAAGGTTGATGCCGTTTTGT TAGGTGCTGTGGGTGGTCCTAAATGGGGTACAGGTAGTGTTAGACCTGAACAAGGTTTACTAAAAATCCGTAAAGAACTT CAATTGTACGCCAACTTAAGACCATGTAACTTTGCATCCGACTCTCTTTTAGACTTATCTCCAATCAAGCCACAATTTGC TAAAGGTACTGACTTCGTTGTTGTCAGAGAATTAGTGGGAGGTATTTACTTTGGTAAGAGAAAGGAAGACGATGGTGATG GTGTCGCTTGGGATAGTGAACAATACACCGTTCCAGAAGTGCAAAGAATCACAAGAATGGCCGCTTTCATGGCCCTACAA CATGAGCCACCATTGCCTATTTGGTCCTTGGATAAAGCTAATGTTTTGGCCTCTTCAAGATTATGGAGAAAAACTGTGGA ACCCAACCCACCTAAATGGTATTATAATCACCAGCAACATGTTTGGTGATATCATCTCCGATGAAGCCTCCGTTATCCCA GGTTCCTTGGGTTTGTTGCCATCTGCGTCCTTGGCCTCTTTGCCAGACAAGAACACCGCATTTGGTTTGTACGAACCATG CCACGGTTCTGCTCCAGATTTGCCAAAGAATAAGGTTGACCCTATCGCCACTATCTTGTCTGCTGCAATGATGTTGAAAT TGTCATTGAACTTGCCTGAAGAAGGTAAGGCCATTGAAGATGCAGTTAAAAAAGGTTTTGGATGCAGGTATCAGAACTGGT GATTTAGGTGGTTCCAACAGTACCACCGAAGTCGGTGATGCTGTCGCCGAAGAAGTTAAGAAAATCCTTGCTTAAAAAAGA TTCTCTTTTTTTATGATATTTGTACATAAACTTTATAAATGAAATTCATAATAGAAACGACACGAAATTACAAAATGGAA CCCGCATGGAATGGGATAATATCACAGGAGGTACTAGACTACCTTTCATCCTACATAAATAGACGCATATAAGTACGCAT CAGTGAGCTGTATGTGCGCAGCTCGCGTTGCATTTTCGGAAGCGCTCGTTTTCGGAAACGCTTTGAAGTTCCTATTCCGA AGTTCCTATTCTCTAGCTAGAAAGTATAGGAACTTCAGAGCGCTTTTGAAAACCAAAAGCGCTCTGAAGACGCACTTTCA ÄAAAACCAAAAACGCACCGGACTGTAACGAGCTACTAAAATATTGCGAATACCGCTTCCACAAACATTGCTCAAAAGTAT CTCTTTGCTATATATCTCTGTGCTATATCCCTATATAACCTACCCATCCACCTTTCGCTCCTTGAACTTGCATCTAAACT CGACCTCTACATTTTTTATGTTTATCTCTAGTATTACTCTTTAGACAAAAAAATTGTAGTAAGAACTATTCATAGAGTGA ATCGAAAACAATACGAAAATGTAAACATTTCCTATACGTAGTATATAGAGACAAAATAGAAGAAAACCGTTCATAATTTTC TGACCAATGAAGAATCATCAACGCTATCACTTTCTGTTCACAAAGTATGCGCAATCCACATCGGTATAGAATATAATCGG GGATGCCTTTATCTTGAAAAAATGCACCCGCAGCTTCGCTAGTAATCAGTAAACGCGGGAAGTGGAGTCAGGCTTTTTTT ATGGAAGAGAAAATAGACACCAAAGTAGCCTTCTTCTAACCTTAACGGACCTACAGTGCAAAAAGTTATCAAGAGACTGC ATTATAGAGCGCACAAAGGAGAAAAAAGTAATCTAAGATGCTTTGTTAGAAAAATAGCGCTCTCGGGATGCATTTTTGT AGAACAAAAAAGAAGTATAGATTCTTTGTTGGTAAAATAGCGCTCTCGCGTTGCATTTCTGTTCTGTAAAAAATGCAGCTC AGATTCTTTGTTTGAAAAATTAGCGCTCTCGCGTTGCATTTTTGTTTTACAAAAATGAAGCACAGATTCTTCGTTGGTAA GCATTTTTGTTCTACAAAATGAAGCACAGATGCTTCGTTAACAAAGATATGCTATTGAAGTGCAAGATGGAAACGCAGAA TCTTCCGTAAAGCGCTAGACTATATTATTATACAGGTTCAAATATACTATCTGTTTCAGGGAAAACTCCCAGGTTCGG ATGTTCAAAATTCAATGATGGGTAACAAGTACGATCCGATATATGCGGTGTGAAATACCGCACAGATGCGTAAGGAGAAA ATACCGCATCAGGCGCCATTCGCCATTCAGGCTGCGCAACTGTTGGGAAGGGCGATCGGTGCGGGCCTCTTCGCTATTAC GCCAGCTGGCGAAAGGGGGATGTGCTGCAAGGCGATTAAGTTGGGTAACGCCAGGGTTTTCCCAGTCACGACGTTGTAAA ACGACGGCCAGTGAATTcatgtaggtggcggaggggagatatacaatagaacagataccagacaagacataatgggctaa acaagactacaccaattacactgcctcattgatggtggtacataacgaactaatactgtagccctagacttgatagccat catcatatcgaagtttcactaccctttttccatttgccatctattgaagtaataataggcgcatgcaacttcttttcttt ttttttcttttctctctcccccgttgttgtctcaccatatccgcaatgacaaaaaatgatggaagacactaaaggaaaa aattaacgacaaagacagcaccaacagatgtcgttgttccagagctgatgaggggtatctcgaagcacacgaaacttttt ccttccttcattgacctgcaattattaatcttttgtttcctcgtcattgttctcgttccctttcttcttttt CTTCTATCGAACAAGCATGCGATATTTGCCGACTTAAAAAGCTCAAGTGCTCCAAAGAAAAACCGAAGTGCGCCAAGTGT CTGAAGAACAACTGGGAGTGTCGCTACTCTCCCAAAACCAAAAGGTCTCCGCTGACTAGGGCACATCTGACAGAAGTGGA ATCAAGGCTAGAAAGACTGGAACAGCTATTTCTACTGATTTTTCCTCGAGAAGACCTTGACATGATTTTGAAAATGGATT GCTTCAGTGGAGACTGATATGCCTCTAACATTGAGACAGCATAGAATAAGTGCGACATCATCATCGGAAGAGAGTAGTAA AATTCCAGCTGACCACCATGTTTatggccagaacaaagcaaacagcaagaaagtccactggtggtaaggccccaagaaag caattagcttctaaggctgccagaaaatccgcccatctaccggtggtgttaagaagcctcacagatataagccaggtac tgttgctttgagagaaatcagaagattccaaaaatctactgaaCCCgggtcccctatactaggttattggaaaggtcgac gcgaccatcctccaaaatcggatctgatcgaaggtcgtggaGATCCCGAAGTTAAACGGGTAAAATTAGAAAACAACGTT GAAGAAATACAACCTGAGCAGGCTGAGACCAATAAACAAGAGGGCACCGATAAAGAGAAATAAAGGAAAGTTCGAGAAAGA AACTGAGAGAATAGGAGGATCTGAAGTGGTTACAGATGTGGAAAAAGGAATTGTCAAATTTGAATTTGATGGTGTTGAAT AAAGAAAACATGATGGTCCTAACTGGATTAAAAAAACATTTTTCAAAAGCAATTACCAAAAAATGCCCAAAGAATACATTGC CAGGTTAGTCTATGATCGAAGTCATCTTTCCATGGCTGTCATTAGGAAGCCATTGACTGTCGTAGGTGGCATAACATATC GACCTTTCGATAAGAGAGAATTCGCAGAAATTGTTTTCTGTGCCATCAGTTCGACGGAACAGGTACGCGGTTATGGTGCG CATCTAATGAATCACTTAAAAGACTATGTTAGAAATACCTCGAACATAAAATATTTTTTGACATATGCAGATAATTACGC ATGAAGGTGGTACGCTGATGCAATGTAACATGGCAATTCCCGGtggcggccgcatcttttacccatacgatgttcctgac tatgcgggctatccctatgacgtcccggactatgcaggatcctatccatatgacgttccagattacgctgctcagtgcgg ccgctctagctagaactagtggatcccccGATACCGTCGACCTGCAGGCATGCAAGCTTGGCGTAATCATGGTCATAGCT GTTTCCTGTGTGAAATTGTTATCCGCTCACAATTCCACACAACATACGAGCCGGAAGCATAAAGTGTAAAGCCTGGGGTG ${\tt CCTAATGAGTGAGCTAACTCACATTAATTGCGTTGCGCTCACTGCCCGCTTTCCAGTCGGGAAACCTGTCGTGCCAGCTG}$ CATTAATGAATCGGCCAACGCGCGGGGAGAGGCGGTTTGCGTATTGGGCGC

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5988-7517: Gal4 DBD-H3-Gcn5-HA

5988-6428: Gal4 DBD (amino acids 1-147)

6501-6587: H4 (amino acids 1-29)

6666-7370: Gcn5 (amino acids 18-252)

7404-7493: trimeric HA

...... 2μ

.....: LEU2
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Translation: Gal4 DBD-H4-Gcn5-HA
MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESRLERLEQLFLLIFPREDLDMI
LKMDSLQDIKALLTGLFVQDNVNKDAVTDRLASVETDMPLTLRQHRISATSSSEESSNKGQRQLTVSNYLFDDEDTPPNP
KKEIEFQLTTMSGRGKGGKGLGKGGAKRHRKILRDNIQGISGSPILGYWKGRRDHPPKSDLIEGRGDPEVKRVKLENNVE
EIQPEQAETNKQEGTDKENKGKFEKETERIGGSEVVTDVEKGIVKFEFDGVEYTFKERPSVVEENEGKIEFRVVNNDNTK
ENMMVLTGLKNIFQKQLPKMPKEYIARLVYDRSHLSMAVIRKPLTVVGGITYRPFDKREFAEIVFCAISSTEQVRGYGAH
LMNHLKDYVRNTSNIKYFLTYADNYAIGYFKKQGFTKEITLDKSIWMGYIKDYEGGTLMQCSMAIPGGGRIFYPYDVPDY
AGYPYDVPDYAGSYPYDVPDYAAQCGRSS

; DNA sequence pdg7 7795 b.p. complete sequence

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ATACATTCTAATGTCTGCCCCTATGTCTGCCCCTAAGAAGATCGTCGTTTTGCCAGGTGACCACGTTGGTCAAGATCACA TGGTGCTGCTATCGATGCTACAGGTGTCCCACTTCCAGATGAGGCGCTGGAAGCCTCCAAGAAGGTTGATGCCGTTTTGT TAGGTGCTGTGGGTGGTCCTAAATGGGGTACAGGTAGTGTTAGACCTGAACAAGGTTTACTAAAAAATCCGTAAAGAACTT CAATTGTACGCCAACTTAAGACCATGTAACTTTGCATCCGACTCTTTTTAGACTTATCTCCAATCAAGCCACAATTTGC TAAAGGTACTGACTTCGTTGTCAGAGAATTAGTGGGAGGTATTTACTTTGGTAAGAGAAAGGAAGACGATGGTGATG GTGTCGCTTGGGATAGTGAACAATACACCGTTCCAGAAGTGCAAAGAATCACAAGAATGGCCGCTTTCATGGCCCTACAA CATGAGCCACCATTGCCTATTTGGTCCTTGGATAAAGCTAATGTTTTGGCCTCTTCAAGATTATGGAGAAAAACTGTGGA ACCCAACCCACCTAAATGGTATTATAATCACCAGCAACATGTTTGGTGATATCATCTCCGATGAAGCCTCCGTTATCCCA GGTTCCTTGGGTTTGTTGCCATCTGCGTCCTTGGCCTCTTTGCCAGACAAGAACACCGCATTTGGTTTGTACGAACCATG ${\tt CCACGGTTCTGCTCCAGATTTGCCAAAGAATAAGGTTGACCCTATCGCCACTATCTTGTCTGCTGCAATGATGTTGAAAT}$ TGTCATTGAACTTGCCTGAAGAAGGTAAGGCCATTGAAGATGCAGTTAAAAAAGGTTTTGGATGCAGGTATCAGAACTGGT GATTTAGGTGGTTCCAACAGTACCACCGAAGTCGGTGATGCTGTCGCCGAAGAAGTTAAGAAAATCCTTGCTTAAAAAAGA TTCTCTTTTTTTTATGATATTTGTACATAAACTTTATAAATGAAATTCATAATAGAAACGACACGAAATTACAAAATGGAA CCCGCATGGAATGGGATAATATCACAGGAGGTACTAGACTACCTTTCATCCTACATAAATAGACGCATATAAGTACGCAT CAGTGAGCTGTATGTGCGCAGCTCGCGTTGCATTTTCGGAAGCGCTCGTTTTCGGAAACGCTTTGAAGTTCCTATTCCGA AGTTCCTATTCTCTAGCTAGAAAGTATAGGAACTTCAGAGCGCTTTTGAAAAACCAAAAGCGCTCTGAAGACGCACTTTCA AAAAACCAAAAACGCACCGGACTGTAACGAGCTACTAAAATATTGCGAATACCGCTTCCACAAACATTGCTCAAAAGTAT CTCTTTGCTATATATCTCTGTGCTATATCCCTATATAACCTACCCATCCACCTTTCGCTCCTTGAACTTGCATCTAAACT CGACCTCTACATTTTTTATGTTTATCTCTAGTATTACTCTTTAGACAAAAAAATTGTAGTAAGAACTATTCATAGAGTGA ATCGAAAACAATACGAAAATGTAAACATTTCCTATACGTAGTATATAGAGACAAAATAGAAGAAACCGTTCATAATTTTC TGACCAATGAAGAATCATCAACGCTATCACTTTCTGTTCACAAAGTATGCGCAATCCACATCGGTATAGAATATAATCGG GGATGCCTTTATCTTGAAAAAATGCACCCGCAGCTTCGCTAGTAATCAGTAAACGCGGGAAGTGGAGTCAGGCTTTTTTT ATGGAAGAGAAAATAGACACCAAAGTAGCCTTCTTCTAACCTTAACGGACCTACAGTGCAAAAAGTTATCAAGAGACTGC ATTATAGAGCGCACAAAGGAGAAAAAAAGTAATCTAAGATGCTTTGTTAGAAAAAATAGCGCTCTCGGGATGCATTTTTGT AGAACAAAAAAGAAGTATAGATTCTTTGTTGGTAAAATAGCGCTCTCGCGTTGCATTTCTGTTCTGTAAAAATGCAGCTC AGATTCTTTGTTTGAAAAATTAGCGCTCTCGCGTTGCATTTTTGTTTTACAAAAATGAAGCACAGATTCTTCGTTGGTAA GCATTTTTGTTCTACAAAATGAAGCACAGATGCTTCGTTAACAAAGATATGCTATTGAAGTGCAAGATGGAAACGCAGAA TCTTCCGTAAAGCGCTAGACTATATATTATTATACAGGTTCAAATATACTATCTGTTTCAGGGAAAACTCCCAGGTTCGG ATGTTCAAAATTCAATGATGGGTAACAAGTACGATCCGATATATGCGGTGTGAAATACCGCACAGATGCGTAAGGAGAAA ATACCGCATCAGGCGCCATTCGCCATTCAGGCTGCGCAACTGTTGGGAAGGGCGATCGGTGCGGGCCTCTTCGCTATTAC GCCAGCTGGCGAAAGGGGGATGTGCTGCAAGGCGATTAAGTTGGGTAACGCCAGGGTTTTCCCAGTCACGACGTTGTAAA ACGACGGCCAGTGAATTcatgtaggtggcggagggagatatacaatagaacagataccagacaagacataatgggctaa acaagactacaccaattacactgcctcattgatggtggtacataacgaactaatactgtagccctagacttgatagccat catcatatcgaagtttcactaccctttttccatttgccatctattgaagtaataataggcgcatgcaacttcttttcttt ttttttttttttttttctctccccgttgttgtctcaccatatccgcaatgacaaaaaatgatggaagacactaaaggaaaa aattaacgacaaagacagcaccaacagatgtcgttgttccagagctgatgaggggtatctcgaagcacacgaaacttttt ccttccttcattgacctgcaattattaatcttttgtttcctcgtcattgttctcgttccctttcttccttgtttctttt ctqcacaatatttcaaqctataccaaqcatacaatcaactccaaqcttgaaqcaaqcctcctgaaaqATGAAGCTACTGT CTTCTATCGAACAAGCATGCGATATTTGCCGACTTAAAAAGCTCAAGTGCTCCAAAGAAAAACCGAAGTGCGCCAAGTGT CTGAAGAACAACTGGGAGTGTCGCTACTCTCCCAAAACCAAAAGGTCTCCGCTGACTAGGGCACATCTGACAGAAGTGGA ATCAAGGCTAGAAAGACTGGAACAGCTATTTCTACTGATTTTTCCTCGAGAAGACCTTGACATGATTTTGAAAATGGATT GCTTCAGTGGAGACTGATATGCCTCTAACATTGAGACAGCATAGAATAAGTGCGACATCATCATCGGAAGAGAGTAGTAA AATTCCAGCTGACCACCATGTCCGGTAGAGGTAAAGGTGGTAAAGGTCTAGGAAAAGGTGGTGCCAAGCGTCACAGAAAG ATTCTAAGAGATAACATCCAAGGTATTTCCgggtcccctatactaggttattggaaaggtcgacgcgaccatcctccaaa

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5988-7517: Gal4 DBD-H3-Gcn5-HA

5988-6428: Gal4 DBD (amino acids 1-147)

6501-6587: H4 (amino acids 1-29)

6666-7370: Gcn5 (amino acids 18-252)

7275-7277: F221A

7404-7493: trimeric HA

...... 2μ

.....: LEU2
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Translation: Gal4 DBD-H4-Gcn5 F221A-HA
MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESRLERLEQLFLLIFPREDLDMI
LKMDSLQDIKALLTGLFVQDNVNKDAVTDRLASVETDMPLTLRQHRISATSSSEESSNKGQRQLTVSNYLFDDEDTPPNP
KKEIEFQLTTMSGRGKGGKGLGKGGAKRHRKILRDNIQGISGSPILGYWKGRRDHPPKSDLIEGRGDPEVKRVKLENNVE
EIQPEQAETNKQEGTDKENKGKFEKETERIGGSEVVTDVEKGIVKFEFDGVEYTFKERPSVVEENEGKIEFRVVNNDNTK
ENMMVLTGLKNIFQKQLPKMPKEYIARLVYDRSHLSMAVIRKPLTVVGGITYRPFDKREFAEIVFCAISSTEQVRGYGAH
LMNHLKDYVRNTSNIKYFLTYADNYAIGYAKKQGFTKEITLDKSIWMGYIKDYEGGTLMQCSMAIPGGGRIFYPYDVPDY
AGYPYDVPDYAGSYPYDVPDYAAQCGRSS

; DNA sequence pdg7 7795 b.p. complete sequence

AATACGGTTATCCACAGAATCAGGGGATAACGCAGGAAAGAACATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCGTA AAAAGGCCGCGTTGCTGGCGTTTTTCCATAGGCTCCGCCCCCTGACGAGCATCACAAAAATCGACGCTCAAGTCAGAGG TGGCGAAACCCGACAGGACTATAAAGATACCAGGCGTTTCCCCCTGGAAGCTCCCTCGTGCGCTCTCCTGTTCCGACCCT GCCGCTTACCGGATACCTGTCCGCCTTTCTCCCTTCGGGAAGCGTGGCGCTTTCTCATAGCTCACGCTGTAGGTATCTCA ${\tt GTTCGGTGTAGGTCGTTCGCTCCAAGCTGGGCTGTGTGCACGAACCCCCCGTTCAGCCCGACCGCTGCGCCTTATCCGGT}$ **AACTATCGTCTTGAGTCCAACCCGGTAAGACACGACTTATCGCCACTGGCAGCCACTGGTAACAGGATTAGCAGAGC** GAGGTATGTAGGCGGTGCTACAGAGTTCTTGAAGTGGTGGCCTAACTACGGCTACACTAGAAGGACAGTATTTGGTATCT TGACGCTCAGTGGAACGAAAACTCACGTTAAGGGATTTTGGTCATGAGATTATCAAAAAGGATCTTCACCTAGATCCTTT TAAATTAAAAATGAAGTTTAAATCAATCTAAAGTATATATGAGTAAACTTGGTCTGACAGTTACCAATGCTTAATCAGT GAGGCACCTATCTCAGCGATCTGTCTATTTCGTTCATCCATAGTTGCCTGACTCCCCGTCGTGTAGATAACTACGATACG GGAGGGCTTACCATCTGGCCCCAGTGCTGCAATGATACCGCGAGACCCACGCTCACCGGCTCAGATTTATCAGCAATAA GAAGCTAGAGTAAGTTCGCCAGTTAATAGTTTGCGCAACGTTGTTGCCATTGCTACAGGCATCGTGGTGTCACGCTC GTCGTTTGGTATGGCTTCATTCAGCTCCGGTTCCCAACGATCAAGGCGAGTTACATGATCCCCCATGTTGTGCAAAAAAG CATAATTCTCTTACTGTCATGCCATCCGTAAGATGCTTTTCTGTGACTGGTGAGTACTCAACCAAGTCATTCTGAGAATA GTGTATGCGGCGACCGAGTTGCTCTTGCCCGGCGTCAATACGGGATAATACCGCGCCACATAGCAGAACTTTAAAAGTGC TCATCATTGGAAAACGTTCTTCGGGGCGAAAACTCTCAAGGATCTTACCGCTGTTGAGATCCAGTTCGATGTAACCCACT CGTGCACCCAACTGATCTTCAGCATCTTTTACTTTCACCAGCGTTTCTGGGTGAGCAAAAACAGGAAGGCAAAATGCCGC AAAAAAGGGAATAAGGGCGACACGGAAATGTTGAATACTCATACTCTTCCTTTTTCAATATTATTGAAGCATTTATCAGG GTTATTGTCTCATGAGCGGATACATATTTGAATGTATTTAGAAAAAATAAACAAATAGGGGTTCCGCGCACATTTCCCCGA AAAGTGCCACCTGACGTCTAAGAAACCATTATTATCATGACATTAACCTATAAAAATAGGCGTATCACGAGGCCCTTTCG TCTTCAAGAATTAACTGTGGGAATACTCAGGTATCGTAAGATGCAAGAGTTCGAATCTCTTAGCAACCATTATTTTTTTC CTCAACATAACGAGAACACACGGGGGGGGCTATCGCACAGACAAATTCGATGACTGGAAATTTTTTTGTTAATTTCAGAGGT

CGCCGCGCATATACCTTTTTCAACTGAAAAATTGGGAGAAAAAGGAAAGGTGAGAGGCCGGAACCGGCTTTTCATATAGA ATAGAGAAGCGTTCATGACTAAATGCTTGCATCACAATACTTGAAGTTGACAATATTATTTAAGGACCTATTGTTTTTC ATACATTCTAATGTCTGCCCCTATGTCTGCCCCTAAGAAGATCGTCGTTTTTGCCAGGTGACCACGTTGGTCAAGATCACA TGGTGCTGCTATCGATGCTACAGGTGTCCCACTTCCAGATGAGGCGCTGGAAGCCTCCAAGAAGGTTGATGCCGTTTTGT TAGGTGCTGTGGGTGGTCCTAAATGGGGTACAGGTAGTGTTAGACCTGAACAAGGTTTACTAAAAATCCGTAAAGAACTT CAATTGTACGCCAACTTAAGACCATGTAACTTTGCATCCGACTCTCTTTTAGACTTATCTCCAATCAAGCCACAATTTGC TAAAGGTACTGACTTCGTTGTTGTCAGAGAATTAGTGGGAGGTATTTACTTTGGTAAGAGAAAGGAAGACGATGGTGATG GTGTCGCTTGGGATAGTGAACAATACACCGTTCCAGAAGTGCAAAGAATCACAAGAATGGCCGCTTTCATGGCCCTACAA CATGAGCCACCATTGCCTATTTGGTCCTTGGATAAAGCTAATGTTTTGGCCTCTTCAAGATTATGGAGAAAAACTGTGGA ACCCAACCACCTAAATGGTATTATAATCACCAGCAACATGTTTGGTGATATCATCTCCGATGAAGCCTCCGTTATCCCA GGTTCCTTGGGTTTGTTGCCATCTGCGTCCTTGGCCTCTTTGCCAGACAAGAACACCGCATTTGGTTTGTACGAACCATG CCACGGTTCTGCTCCAGATTTGCCAAAGAATAAGGTTGACCCTATCGCCACTATCTTGTCTGCCAATGATGTTGAAAT TGTCATTGAACTTGCCTGAAGAAGGTAAGGCCATTGAAGATGCAGTTAAAAAGGTTTTGGATGCAGGTATCAGAACTGGT GATTTAGGTGGTTCCAACAGTACCACCGAAGTCGGTGATGCTGTCGCCGAAGAAGTTAAGAAAATCCTTGCTTAAAAAGA TTCTCTTTTTTTATGATATTTGTACATAAACTTTATAAATGAAATTCATAATAGAAACGACACGAAATTACAAAATGGAA CCCGCATGGAATGGGATAATATCACAGGAGGTACTAGACTACCTTTCATCCTACATAAATAGACGCATATAAGTACGCAT CAGTGAGCTGTATGTGCGCAGCTCGCGTTGCATTTTCGGAAGCGCTCGTTTTCGGAAACGCTTTGAAGTTCCTATTCCGA AGTTCCTATTCTCTAGCTAGAAAGTATAGGAACTTCAGAGCGCTTTTGAAAACCAAAAGCGCTCTGAAGACGCACTTTCA AAAAACCAAAAACGCACCGGACTGTAACGAGCTACTAAAATATTGCGAATACCGCTTCCACAAACATTGCTCAAAAGTAT CTCTTTGCTATATATCTCTGTGCTATATCCCTATATAACCTACCCATCCACCTTTCGCTCCTTGAACTTGCATCTAAACT CGACCTCTACATTTTTTATGTTTATCTCTAGTATTACTCTTTAGACAAAAAAATTGTAGTAAGAACTATTCATAGAGTGA ATCGAAAACAATACGAAAATGTAAACATTTCCTATACGTAGTATATAGAGACAAAATAGAAGAAACCGTTCATAATTTTC TGACCAATGAAGAATCATCAACGCTATCACTTTCTGTTCACAAAGTATGCGCAATCCACATCGGTATAGAATATAATCGG GGATGCCTTTATCTTGAAAAAATGCACCCGCAGCTTCGCTAGTAATCAGTAAACGCGGGAAGTGGAGTCAGGCTTTTTTT ATGGAAGAGAAAATAGACACCAAAGTAGCCTTCTTCTAACCTTAACGGACCTACAGTGCAAAAAGTTATCAAGAGACTGC ATTATAGAGCGCACAAAGGAGAAAAAAAGTAATCTAAGATGCTTTGTTAGAAAAATAGCGCTCTCGGGATGCATTTTTGT AGAACAAAAAAGAAGTATAGATTCTTTGTTGGTAAAATAGCGCTCTCGCGTTGCATTTCTGTTCTGTAAAAAATGCAGCTC AGATTCTTTGTTTGAAAAATTAGCGCTCTCGCGTTGCATTTTTGTTTTACAAAAATGAAGCACAGATTCTTCGTTGGTAA GCATTTTTGTTCTACAAAATGAAGCACAGATGCTTCGTTAACAAAGATATGCTATTGAAGTGCAAGATGGAAACGCAGAA AATGAACCGGGGATGCGACGTGCAAGATTACCTATGCAATAGATGCAATAGTTTCTCCAGGAACCGAAATACATTG TCTTCCGTAAAGCGCTAGACTATATATTATTATACAGGTTCAAATATACTATCTGTTTCAGGGAAAACTCCCAGGTTCGG ATGTTCAAAATTCAATGATGGGTAACAAGTACGATCCGATATATGCGGTGTGAAATACCGCACAGATGCGTAAGGAGAAA ATACCGCATCAGGCGCCATTCGCCATTCAGGCTGCGCAACTGTTGGGAAGGGCGATCGGTGCGGGCCTCTTCGCTATTAC GCCAGCTGGCGAAAGGGGGATGTGCTACAAGGCGATTAAGTTGGGTAACGCCAGGGTTTTCCCAGTCACGACGTTGTAAA ACGACGGCCAGTGAATTcatgtaggtggcggaggggagatatacaatagaacagataccagacaagacataatgggctaa acaagactacaccaattacactgcctcattgatggtggtacataacgaactaatactgtagccctagacttgatagccat catcatatcgaagtttcactaccctttttccatttgccatctattgaagtaataataggcgcatgcaacttcttttcttt ttttttcttttctctctcccccgttgttgtctcaccatatccgcaatgacaaaaaaatgatggaagacactaaaggaaaa aattaacgacaaagacagcaccaacagatgtcgttgttccagaqctgatgaqgggtatctcgaagcacacgaaacttttt cetteetteattgacetgeaattattaatettttgttteetegteattgttetegtteeetttetteettgtttetttt CTTCTATCGAACAAGCATGCGATATTTGCCGACTTAAAAAGCTCAAGTGCTCCAAAGAAAAACCGAAGTGCCCAAGTGT CTGAAGAACAACTGGGAGTGTCGCTACTCTCCCAAAACCAAAAGGTCTCCGCTGACTAGGGCACATCTGACAGAAGTGGA **ATCAAGGCTAGAAAGACTGGAACAGCTATTTCTACTGATTTTTCCTCGAGAAGACCTTGACATGATTTTGAAAATGGATT** GCTTCAGTGGAGACTGATATGCCTCTAACATTGAGACAGCATAGAATAAGTGCGACATCATCATCGGAAGAGAGTAGTAA AATTCCAGCTGACCACCATGTCCGGTAGAGGTAAAGGTGGTAAAGGTCTAGGAAAAGGTGGTGCCAAGCGTCACAGAAAG ${\tt ATTCTAAGAGATAACATCCAAGGTATTTCCgggtcccctatactaggttattggaaaggtcgacgcgaccatcctccaaa}$ atcggatctgatcgaaggtcgtggaGATCCCGAAGTTAAACGGGTAAAATTAGAAAACAACGTTGAAGAAATACAACCTG AGCAGGCTGAGACCAATAAACAAGAGGGCACCGATAAAGAGAATAAAGGAAAGTTCGAGAAAAGAAACTGAGAGAATAGGA GGATCTGAAGTGGTTACAGATGTGGAAAAAGGAATTGTCAAATTTGAATTTGATGGTGTTGAATACACATTCAAAGAGAG TCCTAACTGGATTAAAAAACATTTTTCAAAAGCAATTACCAAAAATGCCCAAAGAATACATTGCCAGGTTAGTCTATGAT CGAAGTCATCTTTCCATGGCTGTCATTAGGAAGCCATTGACTGTCGTAGGTGGCATAACATATCGACCTTTCGATAAGAG AGAATTCGCAGAAATTGTTTTCTGTGCCATCAGTTCGACGGAACAGGTACGCGGTTATGGTGCGCATCTAATGAATCACT TAAAAGACTATGTTAGAAATACCTCGAACATAAAATATTTTTTGACATATGCAGATAATTACGCTATTGGATACGCTAAA ${\tt GATGCAATGTtccATGGCAATTCCCGGtggcggccgcatcttttacccatacgatgttcctgactatgcgggctatccct}$ atqacqtcccqqactatqcaqqatcctatccatatqacqttccagattacqctqctcaqtgcgqccqctctaqctaqaac tagtggatcccccGATACCGTCGACCTGCAGGCATGCAAGCTTGGCGTAATCATGGTCATAGCTGTTTCCTGTGTGAAAT ACTCACATTAATTGCGTTGCGCTCACTGCCCGCTTTCCAGTCGGGAAACCTGTCGTGCCAGCTGCATTAATGAATCGGCC AACGCGCGGGGAGAGGCGGTTTGCGTATTGGGCGC

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434-1861: Gal4 DBD-H3-Gcn5-HA
434-874: Gal4 DBD (amino acids 1-147)
1010-1714: Gcn5 (amino acids 18-252)
1748-1837: trimeric HA
...-..: CEN/ARS
...-..: TRP1
```

Translation: Gal4-DBD-Gcn5-HA
MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESRLERLEQLFLLIFPREDLDMI
LKMDSLQDIKALLTGLFVQDNVNKDAVTDRLASVETDMPLTLRQHRISATSSSEESSNKGQRQLTVSNYLFDDEDTPPNP
KKEIEFQGSPILGYWKGRRDHPPKSDLIEGRGDPEVKRVKLENNVEEIQPEQAETNKQEGTDKENKGKFEKETERIGGSE
VVTDVEKGIVKFEFDGVEYTFKERPSVVEENEGKIEFRVVNNDNTKENMMVLTGLKNIFQKQLPKMPKEYIARLVYDRSH
LSMAVIRKPLTVVGGITYRPFDKREFAEIVFCAISSTEQVRGYGAHLMNHLKDYVRNTSNIKYFLTYADNYAIGYFKKQG
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; DNA sequence PDG28 8087 b.p. complete sequence

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AACATATCGACCTTTCGATAAGAGAGAATTCGCAGAAATTGTTTTCTGTGCCATCAGTTCGACGGAACAGGTACGCGGTT ATGGTGCGCATCTAATGAATCACTTAAAAGACTATGTTAGAAATACCTCGAACATAAAATATTTTTTGACATATGCAGAT TAAAGATTATGAAGGTGGTACGCTGATGCAATGTAACATGGCAATTCCCGGtggcggccgcatcttttacccatacgatg ttcctgactatgcgggctatccctatgacgtcccggactatgcaggatcctatccatatgacgttccagattacgctgct cagtgcggccgctctagctagaactagtggatcccccGATACCGTCGACCTGCAGAGATCTAtgaatcgtagatactgaa ${\tt tgtaactatactcctctaagtttcaatcttggccatgtaacctctgatctatagaattttttaaatgactagaattaatg}$ $\verb|ccc| atcttttttttggacctaaattcttcatgaaaatatattacgagggcttattcagaagctttggacttcttcgccag|$ aggtttggtcaagtctccaatcaaggttgtcggcttgtctaccttgccagaaatttacgaaaagatggaaaagggtcaaa aaaaaaataagtgtatacaaattttaaagtgactcttaggttttaaaacgaaaattcttattcttgagtaactctttcctgtaggtcaggttgctttctcaggtatagcatgaggtcgctcttattgaccacacctctaccggcatgccgagcaaatgcc ${\tt tgcaaatcgctccccatttcacccaattgtagatatgctaactccagcaatgagttgatgaatctcggtgtattttat}$ gtcctcagaggacaacacctgttgtaatcgttcttccacacggatcctggcgtaatagcgaagaggcccgcaccgatcgc 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agcgaaaagacgataaatacaagaaaatgattacgaggatacggagagggtatgtacatgtgtatttatatactaagct $\tt gccggcggttgtttgcaagaccgagaaaaggctagcaagaatcgggtcattgtagcgtatgcgcctgtgaacattctctt$ TTTGCTATTTTGTTAGAGTCTTTTACACCATTTGTCTCCACACCTCCGCTTACATCAACACCAATAACGCCATTTAATCT ${\tt AAGCGCATCACCAACATTTTCTGGCGTCAGTCCACCAGCTAACATAAAATGTAAGCTCTCGGGGCTCTCTTGCCTTCCAA}$ $\verb|CCCAGTCAGAAATCGAGTTCCAAAAGTTCACCTGTCCCACCTGCTTCTGAATCAAACAAGGGAATAAACGAATGA|\\$ GGTTTCTGTGAAGCTGCACTGAGTATGTTGCAGTCTTTTGGAAATACGAGTCTTTTAATAACTGGCAAACCGAGGAA CTCTTGGTATTCTTGCCACGACTCATCTCCATGCAGTTGGACGATATCAATGCCGTAATCATTGACCAGAGCCAAAACAT CCTCCTTAGGTTGATTACGAAACACGCCAACCAAGTATTTCGGAGTGCCTGAACTATTTTTATATGCTTTTACAAGACTT GAAATTTTCCTTGCAATAACCGGGTCAATTGTTCTCTTTCTATTGGGCACACATATAATACCCAGCAAGTCAGCATCGGA ATCTAGTGCACATTCTGCGGCCTCTGTGCTCTGCAAGCCGCAAACTTTCACCAATGGACCAGAACTACCTGTGAAATTAA ${\tt TAACAGACAT} act c caaget {\tt gcctttgtgtgcttaat} caegtatact {\tt caegtgctcaatagtcaccaatgccctccctct}$ ${\tt tggccctctccttttctttttcgaccgaattaattcttgaagacgaaagggcctcgtgatacgcctattttataggtt}$ a at gtcat gataataatg gtttcttagacgtcaggttggcacttttcggggaaatgtgcgcggaacccctatttgtttatt $\verb|tttcta| a atacattca| a atactgta t ccgctcat | gagaca ataaccctga taa atgcttca ataatattga a aagga aga$ $\verb|gtatgagtattcaacatttccgtgtcgcccttattcccttttttgcggcattttgccttcctgtttttgcccagaa|$ ${\tt ACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTACATCGAACTGGATCTCAACAGCGGTAA}$ ${\tt GATCCTTGAGAGTTTTCGCCCCGAAGAACGTTTTCCAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTATTAT}$ $\verb|CCCGTATTGACGCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACTTGGTTGAGTACTCACCAGTC|\\$ ACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATGCAGTGCTGCCATAACCATGAGTGATAACACTGCGGC CAACTTACTTCTGACAACGATCGGAGGACCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACTCGCC TTGATCGTTGGGAACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGACACCACGATGCCTGTAGCAATGGCAACA ${\tt CTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGCCCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCA}$ ACTATGGATGAACGAAATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATTGGtaactgtcagaccaagttta ctcatatatactttagattgatttaaaacttcatttttaatttaaaaggatctaggtgaagatcctttttgataatctca tgaccaaaatcccttaacgtgagttttcgttccactgagcgtcagaccccgtagaaaagatcaaaggatcttcttgagat gctaccaactctttttccgaaggtaactggcttcagcagagcgcagataccaaatactgtccttctagtgtagccgtagt taggccaccacttcaagaactctgtagcaccgcctacatacctcgctctgctaatcctgttaccagtggctgctgccagt ggcgataagtcgtgtcttaccgggttggactcaagacgatagttaccggataaggcgcagcggtcgggctgaacgggggg ttcgtgcacacagcccagcttggagcgaacgacctacaccgaactgagatacctacagcgtgagctatgagaaagcgcca cgcttcccgaagggagaaaggcggacaggtatccggtaagcggcagggtcggaacaggagagcgcacgagggagcttcca gggggaaacgcctggtatctttatagtcctgtcgggtttcgccacctctgacttgagcgtcgatttttgtgatgctcgtc aggggggggggggcctatggaaaaacgccagcaacgcggcctttttacggttcctggccttttgctggccttttgctcaca cgaacgaccgagcgcagcgagtcagtgagcgagggaagcggaagagcgcccaatacgcaaaccgcctctccccgcgcgttg gccgattcattaatgcaggatccgggatcgaagaaatgatggtaaatgaaatgaaatcaaggagcatgaaggcaaaag 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qqacttcctcttttctqqcaaccaaacccatacatcqqqattcctataataccttcqttqqtctccctaacatqtaqqtq $\tt gcggaggggagatatacaatagaacagataccagacaagacataatgggctaaacaagactacaccaattacactgcctc$ attgatg

11

434-1861: Gal4 DBD-H3-Gcn5-HA
434-874: Gal4 DBD (amino acids 1-147)
1010-1714: Gcn5 (amino acids 18-252)
1619-1621: F221A mutation
1748-1837: trimeric HA
.....: CEN/ARS
.....: TRP1

Translation: Gal4-DBD-Gcn5-HA

MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESRLERLEQLFLLIFPREDLDMI LKMDSLQDIKALLTGLFVQDNVNKDAVTDRLASVETDMPLTLRQHRISATSSSEESSNKGQRQLTVSNYLFDDEDTPPNP KKEIEFQGSPILGYWKGRRDHPPKSDLIEGRGDPEVKRVKLENNVEEIQPEQAETNKQEGTDKENKGKFEKETERIGGSE VVTDVEKGIVKFEFDGVEYTFKERPSVVEENEGKIEFRVVNNDNTKENMMVLTGLKNIFQKQLPKMPKEYIARLVYDRSH LSMAVIRKPLTVVGGITYRPFDKREFAEIVFCAISSTEQVRGYGAHLMNHLKDYVRNTSNIKYFLTYADNYAIGYAKKQG FTKEITLDKSIWMGYIKDYEGGTLMQCNMAIPGGGRIFYPYDVPDYAGYPYDVPDYAGSYPYDVPDYAAQCGRSS

; DNA sequence PDG28 8087 b.p. complete sequence

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TACATTGCCAGGTTAGTCTATGATCGAAGTCATCTTTCCATGGCTGTCATTAGGAAGCCATTGACTGTCGTAGGTGGCAT AACATATCGACCTTTCGATAAGAGAGAATTCGCAGAAATTGTTTTCTGTGCCATCAGTTCGACGGAACAGGTACGCGGTT ATGGTGCGCATCTAATGAATCACTTAAAAGACTATGTTAGAAATACCTCGAACATAAAATATTTTTTGACATATGCAGAT TAAAGATTATGAAGGTGGTACGCTGATGCAATGTAACATGGCAATTCCCGGtggcggccgcatcttttacccatacgatg $\verb|tcctgactatgcgggctatccctatgacgtcccggactatgcaggatcctatccatatgacgttccagattacgctgct|$ cagtgcggccgctctagctagaactagtggatcccccGATACCGTCGACCTGCAGAGATCTAtgaatcgtagatactgaa tgtaactatactcctctaagtttcaatcttggccatgtaacctctgatctatagaattttttaaatgactagaattaatg cccatcttttttttggacctaaattcttcatgaaaatatattacgagggcttattcagaagctttggacttcttcgccag aggtttggtcaagtctccaatcaaggttgtcggcttgtctaccttgccagaaatttacgaaaagatggaaaagggtcaaa aaaaaaataagtgtatacaaattttaaagtgactcttaggttttaaaaacgaaaattcttattcttgagtaactctttcct gtaggtcaggttgctttctcaggtatagcatgaggtcgctcttattgaccacacctctaccggcatgccgagcaaatgcc tgcaaatcgctccccatttcacccaattgtagatatgctaactccagcaatgagttgatgaatctcggtgtgtattttat 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ATCTAGTGCACATTCTGCGGCCTCTGTGCTCTGCAAGCCGCAAACTTTCACCAATGGACCAGAACTACCTGTGAAATTAA ${\tt tggccctctccttttcttttcgaccgaattaattcttgaagacgaaagggcctcgtgatacgcctattttataggtt}$ ${\tt aatgtcatgataataatggtttcttagacgtcaggtggcacttttcggggaaatgtgcgcggaacccctatttgtttatt}$ tttctaaatacattcaaatatgtatccgctcatgagacaataaccctgataaatgcttcaataatattgaaaaaggaaga **GENERAL TRANSPORT OF THE PROPERTY OF THE PROP** ACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTACATCGAACTGGATCTCAACAGCGGTAA ${\tt GATCCTTGAGAGTTTTCGCCCCGAAGAACGTTTTCCAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTATTAT}$ $\tt CCCGTATTGACGCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACTTGGTTGAGTACTCACCAGTC$ ACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATGCAGTGCTGCCATAACCATGAGTGATAACACTGCGGC CAACTTACTTCTGACAACGATCGGAGGACCGAAGGAGCTAACCGCTTTTTTTGCACAACATGGGGGATCATGTAACTCGCC TTGATCGTTGGGAACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGACACCACGATGCCTGTAGCAATGGCAACA CTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGCCCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCA ACTATGGATGAACGAAATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATTGGtaactgtcagaccaagttta $\verb"ctcatatatactttagattgatttaaaacttcattttaatttaaaaaggatctaggtgaagatcctttttgataatctca"$ ${\tt tgacca} a a a a tccctta a cgtgagttttcgttccactgagcgtcagaccccgtagaa a agatca a aggatcttcttgagat$ gctaccaactctttttccgaaggtaactggcttcagcagagcgcagataccaaatactgtccttctagtgtagccgtagt ${\tt taggccaccacttcaagaactctgtagcaccgcctacatacctcgctctgctaatcctgttaccagtggctgctgccagt}$ ggcgataagtcgtgtcttaccgggttggactcaagacgatagttaccggataaggcgcagcggtcgggctgaacggggg ttcgtgcacacagcccagcttggagcgaacgacctacaccgaactgagatacctacagcgtgagctatgagaaagcgcca

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434-1894: Gal4 DBD-MCS-Gcn5(wt)-HA 926-970: multicloning sequence 1043-1747: Gcn5 (aa18-252) 1781-1871: trimeric HA

Translation: GDBD-MCS-Gcn5-HA
MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESRLERLEQLFLLIFPREDLDMI
LKMDSLQDIKALLTGLFVQDNVNKDAVTDRLASVETDMPLTLRQHRISATSSSEESSNKGQRQLTVSNYLFDDEDTPPNP
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EKETERIGGSEVVTDVEKGIVKFEFDGVEYTFKERPSVVEENEGKIEFRVVNNDNTKENMMVLTGLKNIFQKQLPKMPKE
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QCGRSS

; ### from DNA Strider Wednesday, March 26, 2003 1:40:57 PM; DNA sequence pDG30 8120 b.p. complete sequence

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434-1894: Gal4 DBD-MCS-Gcn5(wt)-HA 926-970: multicloning sequence 1043-1747: Gcn5 (F221A) (aal8-252) 1781-1871: trimeric HA

Translation: GDBD-MCS-Gcn5(F221A)-HA

MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESRLERLEQLFLLIFPREDLDMI LKMDSLQDIKALLTGLFVQDNVNKDAVTDRLASVETDMPLTLRQHRISATSSSEESSNKGQRQLTVSNYLFDDEDTPPNP KKEIEFQGTMHELPRLEPGSPILGYWKGRRDHPPKSDLIEGRGDPEVKRVKLENNVEEIQPEQAETNKQEGTDKENKGKF EKETERIGGSEVVTDVEKGIVKFEFDGVEYTFKERPSVVEENEGKIEFRVVNNDNTKENMMVLTGLKNIFQKQLPKMPKE YIARLVYDRSHLSMAVIRKPLTVVGGITYRPFDKREFAEIVFCAISSTEQVRGYGAHLMNHLKDYVRNTSNIKYFLTYAD NYAIGYAKKQGFTKEITLDKSIWMGYIKDYEGGTLMQCNMAIPGGGRIFYPYDVPDYAGYPYDVPDYAGSYPYDVPDYAA QCGRSS

; ### from DNA Strider Wednesday, March 26, 2003 1:41:20 PM; DNA sequence pDG31 8120 b.p. complete sequence

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Translation: Ga14-DBD-p53(300-393)-Gcn5-HA

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434-2146: Gal4 DBD-p53-Gcn5-HA

434-874: Gal4 DBD (amino acids 1-147)

938-1216: p53 (amino acids 300-393)

1295-1999: Gcn5 (amino acids 18-252) with Phe221 changed to Ala

2033-2119: trimeric HA
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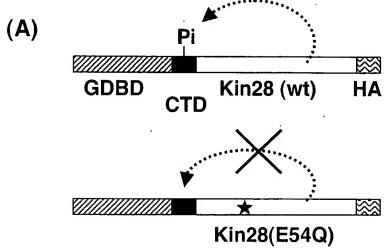
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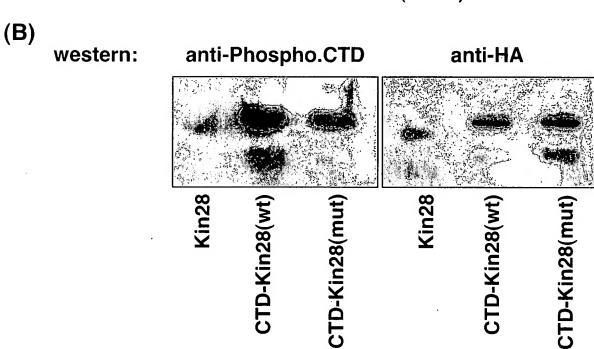
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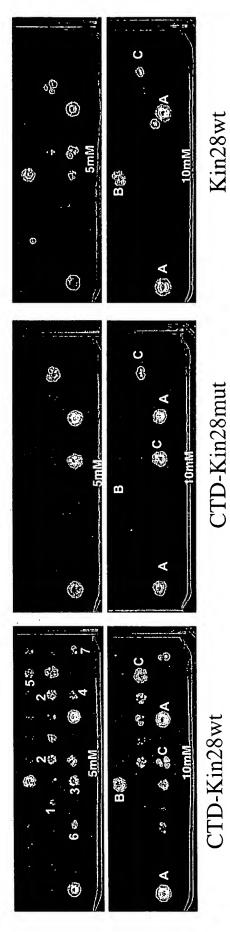
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 $\verb|ccgATACCGTCGACCTGCAGAGATCTAtgaatcgtagatactgaaaaaccccgcaagttcacttcaactgtgcatcgtgc|\\$ accatctcaatttctttcatttatacatcgttttgccttcttttatgtaactatactcctctaagtttcaatcttggcca at a tatattac gagggcttattcaga agctttggacttcttcgccagaggtttggtcaagtctccaatcaaggttgtcggct ${\tt tgtctaccttgccagaaatttacgaaaagatggaaaagggtcaaatcgttggtagatacgttgttgacacttctaaataa}$ $\verb|ttaggttttaaaacgaaaattcttattcttgagtaactctttcctgtaggtcaggttgctttctcaggtatagcatgagg|$ $\verb|tcgctcttattgaccacctctaccggcatgccgagcaaatgcctgcaaatcgctccccatttcacccaattgtagata|$ ${\tt tgctaactccagcaatgagttgatgaatctcggtgtgtattttatgtcctcagaggacaacacctgttgtaatcgttctt}$ $\verb|ccacacggatcctggcgta| tagcgaagaggcccgcaccgatcgccttcccaacagttgcgcagcctgaatggcgaatg$ $\tt gcgcctgatgcggtatttctccttacgcatctgtgcggtatttcacaccgcatatatcgctgggccattctcatgaaga$ atatettgaatttattgteatattaetagttggtgtggaagteeatatateggtgateaatatagtggttgaeatgetgg ctagtcaacattgagccttttgatcatgcaaatatattacggtattttacaatcaaatatcaaacttaactattgacttt ataacttatttaggtggtaacattcttataaaaaagaaaaaattactgcaaaacagtactagcttttaacttgtatcct aatacacaaatctggcttaataaagtctataatatctcataaagaagtgctaaattggctagtgctatatatttttaa gaaaatttettttgactaagteeatategactttgtaaaagtteactttagcatacatatattacaegageeagaaattg cctgctcaagaaaaagaaactgttttgtccttggaaaaaaagcactacctaggagcggccaaaatgccgaggctttcata tgaaggagcatgttcggcacacagtggaccgaacgtggggtaagtgcactagggtccggttaaacggatctcgcattgat tggtgtttatgcaaagaaaccactgtgtttaatatgtgtcactgtttgatattactgtcagcgtagaagataatagtaaa agcggttaataagtgtatttgagataagtgtgataaagtttttacagcgaaaagacgataaatacaagaaaatgattacg ${\tt aggatacggagagaggtatgtacatgtgtatttatatactaagctgccggcggttgtttgcaagaccgagaaaaggctag}$ ${\tt caagaatcgggtcattgtagcgtatgcgcctgtgaacattctcttcaacaagtttgattccattgcggtgaaatggtaaa}$ ${ t aaatactactcagtaataacctaTTTCTTAGCATTTTTGACGAAATTTGCTATTTTGTTAGAGTCTTTTACACCATTTGT}$ CTCCACACCTCCGCTTACATCAACACCAATAACGCCATTTAATCTAAGCGCATCACCAACATTTTCTGGCGTCAGTCCAC CAGCTAACATAAAATGTAAGCTCTCGGGGGCTCTCTTGCCTTCCAACCCAGTCAGAAATCGAGTTCCAATCCAAAAGTTCA CCTGTCCCACCTGCTTCTGAATCAAACAAGGGAATAAACGAATGAGGTTTCTGTGAAGCTGCACTGAGTAGTATGTTGCA GTCTTTTGGAAATACGAGTCTTTTAATAACTGGCAAACCGAGGAACTCTTGGTATTCTTGCCACGACTCATCTCCATGCA TATTTCGGAGTGCCTGAACTATTTTATATGCTTTTACAAGACTTGAAATTTTCCTTGCAATAACCGGGTCAATTGTTCT $\verb|CTTTCTATTGGGCACACATATAATACCCAGCAAGTCAGCATCGGAATCTAGTGCACATTCTGCGGCCTCTGTGCTCTGCA| \\$ ${\tt AGCCGCAAACTTTCACCAATGGACCAGAACTACCTGTGAAATTAATAACAGACATactccaagctgcctttgtgtgctta}$ tcttgaagacgaaagggcctcgtgatacgcctatttttataggttaatgtcatgataataatggtttcttagacgtcagg tggcacttttcgggggaaatgtgcgcggaacccctatttgtttatttttctaaatacattcaaatatgtatccgctcatga gacaataaccctgataaatgcttcaataatattgaaaaaggaagagtATGAGTATTCAACATTTCCGTGTCGCCCTTATT CCCTTTTTTGCGGCATTTTGCCTTCCTGTTTTTGCTCACCCAGAAACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTT GGGTGCACGAGTGGGTTACATCGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTTTTCGCCCCGAAGAACGTTTTC CAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTATTATCCCGTATTGACGCCGGGCAAGAGCAACTCGGTCGC CGCATACACTATTCTCAGAATGACTTGGTTGAGTACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAG $\verb|CCAAACGACGAGGCGTGACACCACGATGCCTGTAGCAATGGCAACAACGTTGCGCAAACTATTAACTGGCGAACTACTTAC|$ TCTAGCTTCCCGGCAACAATTAATAGACTGGATGGAGGCGGATAAAGTTGCAGGACCACTTCTGCGCTCGGCCCTTCCGG ${\tt AAGCCCTCCCGTATCGTAGTTATCTACACGACGGGGGGGTCAGGCAACTATGGATGAACGAAATAGACAGATCGCTGAGAT}$







Phosphorylated CTD-interacting proteins

Kin28-interacting proteins

C = YDL100c

B = Pc110

A = Cc11

1 = Fcp12 = Ssn8/Srb11

3= Tfb3

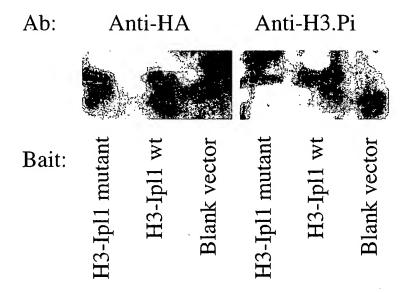
4 = Whi2

5 = YMR181c6 = YPL229w

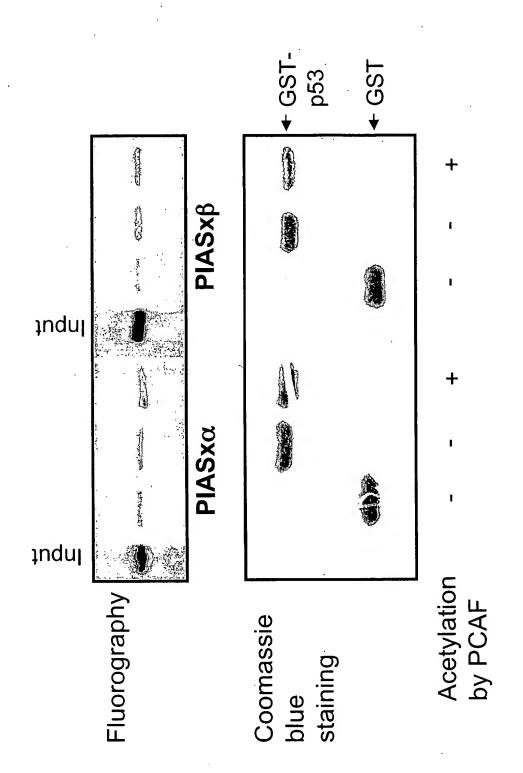
7 = YDR428C

FIG. 27

Autophosphorylation of H3 at Ser10 by the tethered Ipl1 kinase



PIASx α and acetylation-independent interaction between p53 and PIASx β GST pulldown shows acetylation-stimulated interaction between p53 and



434-2482: Gal4 DBD-H3-Ipl1-HA

434-874: Gal4 DBD (amino acids 1-147) 932-952: TEV cleavage sequence (ENLYFQG)

974-1153: H3 (amino acids 1-59)

1235-2332: Ipl1 ORF

2369-2458: HAx3

Translation: Gal4 DBD-H3-Ipl1-Hax3

MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESR
LERLEQLFLLIFPREDLDMILKMDSLQDIKALLTGLFVQDNVNKDAVTDRLASVETDMPL
TLRQHRISATSSSEESSNKGQRQLTVSNYLFDDEDTPPNPKKEIEFQENLYFQGLTTMFM
ARTKQTARKSTGGKAPRKQLASKAARKSAPSTGGVKKPHRYKPGTVALREIRRFQKSTEP
GSPILGYWKGRRDHPPKSDLIEGRGGPQRNSLVNIKLNANSPSKKTTTRPNTSRINKPWR
ISHSPQQRNPNSKIPSPVREKLNRLPVNNKKFLDMESSKIPSPIRKATSSKMIHENKKLP
KFKSLSLDDFELGKKLGKGKFGKVYCVRHRSTGYICALKVMEKEEIIKYNLQKQFRREVE
IQTSLNHPNLTKSYGYFHDEKRVYLLMEYLVNGEMYKLLRLHGPFNDILASDYIYQIANA
LDYMHKKNIIHRDIKPENILIGFNNVIKLTDFGWSIINPPENRRKTVCGTIDYLSPEMVE
SREYDHTIDAWALGVLAFELLTGAPPFEEEMKDTTYKRIAALDIKMPSNISQDAQDLILK
LLKYDPKDRMRLGDVKMHPWILRNKPFWENKRLELMAIPGGGRIFYPYDVPDYAGYPYDV
PDYAGSYPYDVPDYAAQCGRSS

; ### from DNA Strider Thursday, February 5, 2004 12:57:23 PM ; DNA sequence PDG64 8708 b.p. complete sequence

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434-2482: Gal4 DBD-H3-Ipl1-HA 434-874: Gal4 DBD (amino acids 1-147) 932-952: TEV cleavage sequence (ENLYFQG) 974-1153: H3 (amino acids 1-59) 1235-2332: Ipl1 ORF with E152Q V153L double mutations 2369-2458: HAx3

Translation: Gal4 DBD-H3-Ipl1-HAx3

MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESR

LERLEQLFLLIFPREDLDMILKMDSLQDIKALLTGLFVQDNVNKDAVTDRLASVETDMPL

TLRQHRISATSSSEESSNKGQRQLTVSNYLFDDEDTPPNPKKEIEFQENLYFQGLTTMFM

ARTKQTARKSTGGKAPRKQLASKAARKSAPSTGGVKKPHRYKPGTVALREIRRFQKSTEP

GSPILGYWKGRRDHPPKSDLIEGRGGPQRNSLVNIKLNANSPSKKTTTRPNTSRINKPWR

ISHSPQQRNPNSKIPSPVREKLNRLPVNNKKFLDMESSKIPSPIRKATSSKMIHENKKLP

KFKSLSLDDFELGKKLGKGKFGKVYCVRHRSTGYICALKVMEKEEIIKYNLQKQFRRQLE

IQTSLNHPNLTKSYGYFHDEKRVYLLMEYLVNGEMYKLLRLHGPFNDILASDYIYQIANA

LDYMHKKNIIHRDIKPENILIGFNNVIKLTDFGWSIINPPENRRKTVCGTIDYLSPEMVE

SREYDHTIDAWALGVLAFELLTGAPPFEEEMKDTTYKRIAALDIKMPSNISQDAQDLILK

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; ### from DNA Strider Thursday, February 5, 2004 12:57:23 PM ; DNA sequence PDG65 8708 b.p. complete sequence

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CGAAAAAGACCACAACAAGACCAAATACGTCCAGGATCAATAAACCATGGAGAATATCCCATTC GCCGCAGCAAAGAAACCCGAATTCAAAAATACCTTCACCTGTAAGAGAAAAATTGAACAGATTA CCTGTAAACAATAAGAAGTTTTTGGATATGGAAAGCTCCAAAATTCCATCACCTATAAGGAAAG CGACTTCTTCCAAAATGATACACGAAAATAAGAAGCTACCTAAATTTAAATCCCTATCACTCGA TGACTTTGAACTGGGGAAGAATTAGGAAAGGGTAAATTCGGTAAAGTTTATTGCGTTCGGCAC ATCATACGGCTATTTTCATGATGAAAAAAGAGTGTACCTGCTAATGGAATACTTAGTCAATGGG GAAATGTATAAACTATTGAGGTTACACGGACCCTTCAACGATATTTTAGCATCAGATTATATTT ATCAAATTGCCAATGCCCTAGATTATATGCATAAAAAGAATATTATTCATAGAGATATTAAACC TGAAAATATACTAATAGGGTTCAATAATGTCATTAAGTTAACGGACTTCGGATGGAGTATAATA AATCCGCCAGAAATAGAAGGAAAACTGTCTGTGGGACAATTGACTACCTTTCTCCAGAAATGG TGGAGTCAAGGGAATATGATCACACTATAGATGCATGGGCTCTTGGCGTCCTGGCGTTTGAACT ACTGACCGGTGCCCCTCCGTTCGAAGAAGAATGAAAGATACTACATATAAAAGGATAGCAGCA CTGGATATCAAAATGCCCAGTAACATTTCTCAGGATGCGCAAGATTTAATACTTAAACTACTAA AATACGACCCCAAAGATAGAATGCGCCTTGGAGACGTAAAAATGCATCCTTGGATACTAAGAAA CAAGCCCTTTTGGGAAAATAAGCGGTTAGAGCTCATGGCAATTCCCGGtqqcqccqcatcttt tacccatacqatqttcctqactatgcqqqctatccctatgacqtcccggactatgcaggatcct atccatatgacqttccagattacqctqctcagtqcgqccqctctagctagaactagtggatccc ccGATACCGTCGACCTGCAGAGATCTAtqaatcqtaqatactqaaaaaccccgcaagttcactt caactgtgcatcgtgcaccatctcaatttctttcatttatacatcgttttgccttcttttatgt aactatactcctctaagtttcaatcttggccatgtaacctctgatctatagaattttttaaatg actagaattaatgcccatcttttttttggacctaaattcttcatgaaaatatattacgagggct tattcagaagctttggacttcttcgccagaggtttggtcaagtctccaatcaaggttgtcggct tqtctaccttqccaqaaatttacqaaaaqatqqaaaaqqgtcaaatcgttggtagatacgttgt tgacacttctaaataagcqaatttcttatgatttatgatttttattattaaataagttataaaa aaaataaqtqtatacaaattttaaaqtqactcttaqgttttaaaacgaaaattcttattcttga gtaactctttcctgtaggtcaggttgctttctcaggtatagcatgaggtcgctcttattgacca cacctctaccggcatgccgagcaaatgcctgcaaatcgctccccatttcacccaattgtagata tgctaactccagcaatgagttgatgaatctcggtgtgtattttatgtcctcagaggacaacacc tgttgtaatcqttcttccacacqqatcctqqcqtaataqcqaaqaqqcccqcaccqatcqccct teccaacagttgegeageetgaatggegaatggegeetgatgeggtatttteteettaegeate tgtgcggtatttcacaccgcatatatcgctgggccattctcatgaagaatatcttgaatttatt gtcatattactagttggtgtggaagtccatatatcggtgatcaatatagtggttgacatgctgg ctagtcaacattgagccttttgatcatgcaaatatattacggtattttacaatcaaatatcaaa cttaactattgactttataacttatttaggtggtaacattcttataaaaaaagaaaaaattact gcaaaacagtactagcttttaacttgtatcctaggttatctatgctgtctcaccatagagaata ttacctatttcagaatgtatgtccatgattcgccgggtaaatacatataatacacaaatctggc ttaataaagtctataatatctcataaagaagtgctaaattggctagtgctatatatttttaa qaaaatttcttttqactaaqtccatatcqactttgtaaaagttcactttagcatacatatatta cataaatqaaataatttatttattqtttatgattaccgaaacataaaacctgctcaagaaaaag aaactqttttqtccttqqaaaaaaqcactacctaggagcggccaaaatgccgaggctttcata gttttatcgtcacagttttacagtaaataagtatcacctcttagagttcgatgataagctgtca aacatgagaattaattccacatgttaaaatagtgaaggagcatgttcggcacacagtggaccga

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acaccgaactgagatacctacagcgtgagctatgagaaagcgccacgcttcccgaagggagaaa ggcggacaggtatccggtaagcggcagggtcggaacaggagcgcacgagggagcttccaggg ggaaacgcctggtatctttatagtcctgtcgggtttcgccacctctgacttgagcgtcgatttt ${\tt tgtgatgctcgtcagggggggggggggcctatggaaaaacgccagcaacgcggcctttttacggtt}$ $\verb|cctggccttttgctcacatgttctttcctgcgttatcccctgattctgtggat|\\$ aaccgtattaccgcctttgagtgagctgataccgctcgccgcagccgaacgaccgagcgcagcg ${\tt agtcagtgagcgaggaagcggcagaagcgcccaatacgcaaaccgcctctccccgcgcgttggcc}$ gattcattaatgcaggatccgggatcgaagaaatgatggtaaatgaaataggaaatcaaggagc $\verb|atgaaggcaaaagacaaatataagggtcgaacgaaaaataaagtgaaaagtgttgatatgatgt|$ $\verb|atttggctttgcggcgcgaaaaaacgagtttacgcaattgcacaatcatgctgactctgtggc|$ ggacccgcgctcttgccggcccggcgataacgctgggcgtgaggctgtgcccggcggagttttt tgcgcctgcattttccaaggtttaccctgcgctaaggggcgagattggagaagcaataagaatg $\verb|ccggttggggttgcgatgatgacgaccacgacaactggtgtcattatttaagttgccgaaagaa|\\$ cctgagtgcatttgcaacatgagtatactagaagaatgagccaagacttgcgagacgcgagttt gccggtggtgcgaacaatagagcgaccatgaccttgaaggtgagacgcgcataaccgctagagt gaaatggttgtctgtttgagtacgctttcaattcatttgggtgtgcactttattatgttacaat $\verb|tccttttgttgtttccgggtgtacaatatggacttcctcttttctggcaaccaaacccatacat|\\$ $\verb|cgggattcctataataccttcgttggtctccctaacatgtaggtggcggaggggagatatacaa|\\$ tagaacagataccagacaagacataatgggctaaacaagactacaccaattacactgcctcattgatg //

434-2041: Gal4 DBD-TEV-Kin28-HAx3 434-874: Gal4 DBD (amino acids 1-147) 932-952: TEV cleavage sequence (ENLYFQG) 953-958: EcoRI 959-964: PuvII 969-974: NcoI 977-1894: Kin28 (cDNA) 1991-2080: trimeric HA

Translation: Gal4 DBD-CTDx3-TEV-Kin28-HAx3

MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESR
LERLEQLFLLIFPREDLDMILKMDSLQDIKALLTGLFVQDNVNKDAVTDRLASVETDMPL
TLRQHRISATSSSEESSNKGQRQLTVSNYLFDDEDTPPNPKKEIELENLYFQGEFQLTTMA
MKVNMEYTKEKKVGEGTYAVVYLGCQHSTGRKIAIKEIKTSEFKDGLDMSAIREVKYLQE
MQHPNVIELIDIFMAYDNLNLVLEFLPTDLEVVIKDKSILFTPADIKAWMLMTLRGVYHC
HRNFILHRDLKPNNLLFSPDGQIKVADFGLARAIPAPHEILTSNVVTRWYRAPELLFGAK
HYTSAIDIWSVGVIFAELMLRIPYLPGQNDVDQMEVTFRALGTPTDRDWPEVSSFMTYNK
LQIYPPPSRDELRKRFIAASEYALDFMCGMLTMNPQKRWTAVQCLESDYFKELPPPSDPS
SIKIRNVMAIPGGGRIFYPYDVPDYAGYPYDVPDYAGSYPYDVPDYAAQCGRSS-N-WIP
RYRRPAEIYES-ILKNPASSLQLCIVHHLNFFHLYIVLPSFM-LYSSKFQSWPCNL-SIE
FFK-LELMPIFFLDLNS

pMK498 w/ TEV 8264 b.p. complete sequence ; DNA sequence gtggtacataacgaactaatactgtagccctagacttgatagccatcatcatatcgaagtttca ctaccctttttccatttgccatctattgaagtaataataggcgcatgcaacttcttttttt ttttcttttctctctcccccgttgttgtctcaccatatccgcaatgacaaaaaatgatggaag acactaaaggaaaaattaacgacaaagacagcaccaacagatgtcgttgttccagagctgatg aggggtatctcgaagcacacgaaactttttccttccttcattgacctgcaattattaatctttt gtttcctcgtcattgttctcgttccctttcttccttgtttcttttttctgcacaatatttcaagc tataccaagcatacaatcaactccaagcttgaagcaagcctcctgaaagATGAAGCTACTGTCT TCTATCGAACAAGCATGCGATATTTGCCGACTTAAAAAGCTCAAGTGCTCCAAAGAAAAACCGA AGTGCGCCAAGTGTCTGAAGAACAACTGGGAGTGTCGCTACTCTCCCAAAACCAAAAGGTCTCC GCTGACTAGGGCACATCTGACAGAAGTGGAATCAAGGCTAGAAAGACTGGAACAGCTATTTCTA CTGATTTTTCCTCGAGAAGACCTTGACATGATTTTGAAAATGGATTCTTTACAGGATATAAAAG TTCAGTGGAGACTGATATGCCTCTAACATTGAGACAGCATAGAATAAGTGCGACATCATCATCG ATACCCCACCAAACCCCAAAAAAAAGAGATCGAATTggagaatttgtattttcagggcgaattcCA

 GTAGCAGATTTCGGTCTAGCAAGGGCGATACCGGCCCCACATGAGATACTGACAAGTAACGTCG TAACAAGATGGTATAGAGCGCCAGAATTGTTGTTTGGAGCTAAACATTACACATCGGCTATTGA TATCTGGTCAGTAGGCGTTATATTCGCGGAATTAATGCTAAGGATACCTTATTTACCAGGACAG AATGATGTCGATCAAATGGAAGTAACGTTCAGGGCCTTAGGGACACCTACAGATAGAGATTGGC CCGAAGTTTCTTCCTTTATGACGTATAACAAGTTACAAATATATCCGCCCCCTTCAAGAGATGA ATTGAGGAAAAGGTTCATTGCTGCTAGCGAATACGCCTTAGATTTTATGTGTGGAATGCTAACG ${\tt ATGAACCCACAAAAGAGTGACCGCTGTTCAGTGTTTAGAAAGTGATTATTTCAAAGAATTAC}$ CACCACCAAGTGACCCGTCTTCAATAAAAATACGTAACgtcatggCAATTCCCGGtggcggccg $\verb"catcttttacccatacgatgttcctgactatgcgggctatccctatgacgtcccggactatgca"$ ggatcccccgataCCGTCGACCTGCAGAGATCTAtgaatcgtagatactgaaaaaccccgcaag $\verb|tttatgtaactatactcctctaagtttcaatcttggccatgtaacctctgatctatagaatttt|\\$ ttaaatgactagaattaatgcccatctttttttttggacctaaattcttcatgaaaatatattac gagggcttattcagaagctttggacttcttcgccagaggtttggtcaagtctccaatcaaggtt $\tt gtcggcttgtctaccttgccagaaatttacgaaaagatggaaaagggtcaaatcgttggtagat$ acgttgttgacacttctaaataagcgaatttcttatgatttatgatttttattattaataagttataaaaaaaataagtgtatacaaattttaaagtgactcttaggttttaaaacgaaaattctta ttcttgagtaactctttcctgtaggtcaggttgctttctcaggtatagcatgaggtcgctctta ttgaccacacctctaccggcatgccgagcaaatgcctgcaaatcgctccccatttcacccaatt $\tt gtagatatgctaactccagcaatgagttgatgaatctcggtgtgtattttatgtcctcagagga$ caacacctgttgtaatcgttcttccacacggatcctggcgtaatagcgaagaggcccgcaccga tcgcccttcccaacagttgcgcagcctgaatggcgaatggcgcctgatgcggtattttctcctt acgcatctgtgcggtatttcacaccgcatatatcgctgggccattctcatgaagaatatcttga atttattgtcatattactagttggtgtggaagtccatatatcggtgatcaatatagtggttgac atgctggctagtcaacattgagccttttgatcatgcaaatatattacggtattttacaatcaaa tatcaaacttaactattgactttataacttatttaggtggtaacattcttataaaaaagaaaaa aattactgcaaaacagtactagcttttaacttgtatcctaggttatctatgctgtctcaccata gagaatattacctatttcagaatgtatgtccatgattcgccgggtaaatacatataatacacaa atctggcttaataaagtctataatatctcataaagaagtgctaaattggctagtgctatata tttttaagaaaatttcttttgactaagtccatatcgactttgtaaaagttcactttagcataca tatattacacgagccagaaattgtaacttttgcctaaaatcacaaattgcaaaatttaattgct ttttaaacataaatgaaataatttatttattgtttatgattaccgaaacataaaacctgctcaa gaaaaagaaactgttttgtccttggaaaaaaagcactacctaggagcggccaaaatgccgaggc tttcatagcttaaactctttacagaaaataggcattatagatcagttcgagttttcttattctt ccttccggttttatcgtcacagttttacagtaaataagtatcacctcttagagttcgatgataa gctgtcaaacatgagaattaattccacatgttaaaatagtgaaggagcatgttcggcacacagt ggaccgaacgtggggtaagtgcactagggtccggttaaacggatctcgcattgatgaggcaacg ctaattatcaacatatagattgttatctatctgcatgaacacgaaatctttacttgacgacttg aggctgatggtgtttatgcaaagaaaccactgtgtttaatatgtgtcactgtttgatattactg tcagcgtagaagataatagtaaaagcggttaataagtgtatttgagataagtgtgataaagttt ttacagcgaaaagacgataaatacaagaaaatgattacgaggatacggagagaggtatgtacat gtgtatttatatactaagctgccggcggttgtttgcaagaccgagaaaaggctagcaagaatcg ggtcattgtagcgtatgcgcctgtgaacattctcttcaacaagtttgattccattgcggtgaaa

TCTTAGCATTTTTGACGAAATTTGCTATTTTGTTAGAGTCTTTTACACCATTTGTCTCCACACC TCCGCTTACATCAACACCAATAACGCCATTTAATCTAAGCGCATCACCAACATTTTCTGGCGTC AGTCCACCAGCTAACATAAAATGTAAGCTCTCGGGGCTCTCTTGCCTTCCAACCCAGTCAGAAA TCGAGTTCCAAAAGTTCACCTGTCCCACCTGCTTCTGAATCAAACAAGGGAATAAACGA ATGAGGTTTCTGTGAAGCTGCACTGAGTAGTATGTTGCAGTCTTTTGGAAATACGAGTCTTTTA ATAACTGGCAAACCGAGGAACTCTTGGTATTCTTGCCACGACTCATCTCCATGCAGTTGGACGA TATCAATGCCGTAATCATTGACCAGAGCCAAAACATCCTCCTTAGGTTGATTACGAAACACGCC AACCAAGTATTTCGGAGTGCCTGAACTATTTTTATATGCTTTTACAAGACTTGAAATTTTCCTT GCAATAACCGGGTCAATTGTTCTCTTTCTATTGGGCACACATATAATACCCAGCAAGTCAGCAT $\tt CGGAATCTAGTGCACATTCTGCGGCCTCTGTGCTCTGCAAGCCGCAAACTTTCACCAATGGACC$ AGAACTACCTGTGAAATTAATAACAGACATactccaagctgcctttgtgtgcttaatcacgtat aattaattcttgaagacgaaagggcctcgtgatacgcctatttttataggttaatgtcatgata ataatggtttcttagacgtcaggtggcacttttcggggaaatgtgcgcggaacccctatttgtt tatttttctaaatacattcaaatatgtatccgctcatgagacaataaccctgataaatgcttca $\verb|ataatattgaaaaaggaagagtATGAGTATTCAACATTTCCGTGTCGCCCTTATTCCCTTTTTT|$ GCGGCATTTTGCCTTCTTTTTGCTCACCCAGAAACGCTGGTGAAAGTAAAAGATGCTGAAG ATCAGTTGGGTGCACGAGTGGGTTACATCGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAG TTTTCGCCCCGAAGAACGTTTTCCAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTA TTATCCCGTATTGACGCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACT TGGTTGAGTACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATG CCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACTCGCCTTGATCGTTGGG AACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGACACCACGATGCCTGTAGCAATGGC TTATTGCTGATAAATCTGGAGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCC AGATGGTAAGCCCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCAACTATGGATGAA CGAAATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATTGGtaactgtcagaccaag tttactcatatatactttagattgatttaaaacttcatttttaatttaaaaggatctaggtgaa gatcctttttgataatctcatgaccaaaatcccttaacgtgagttttcgttccactgagcgtca ttttccgaaggtaactggcttcagcagagcgcagataccaaatactgtccttctagtgtagccg tagttaggccaccacttcaagaactctgtagcaccgcctacatacctcgctctgctaatcctgt taccagtggctgctgccagtggcgataagtcgtgtcttaccgggttggactcaagacgatagtt accggataaggcgcagcggtcggggctgaacggggggttcgtgcacacagcccagcttggagcga acgacctacaccgaactgagatacctacagcgtgagctatgagaaagcgccacgcttcccgaag ggagaaaggcggacaggtatccggtaagcggcagggtcggaacaggagcgcacgagggagct tccagggggaaacgcctggtatctttatagtcctgtcgggtttcgccacctctgacttgagcgt cgatttttgtgatgctcgtcagggggggggggcctatggaaaaacgccagcaacgcggcctttt tacggttcctggccttttgctggccttttgctcacatgttctttcctgcgttatcccctgattc tgtggataaccgtattaccgcctttgagtgagctgataccgctcgccgcagccgaacgaccgag cgcagcgagtcagtgagcgaggaagcggaagagcgccaatacgcaaaccgcctctccccgcgc gttggccgattcattaatgcaggatccgggatcgaagaaatgatggtaaatgaaataggaaatc

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434-2104: Gal4 DBD-TEV-CTDx3-Kin28-HAx3
434-874: Gal4 DBD (amino acids 1-147)
932-952: TEV cleavage sequence (ENLYFQG)
962-1024: CTD (three tandem repeats of YSTPSPS)
1040-1957: Kin28 (cDNA)
1991-2080: trimeric HA
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Translation: Gal4 DBD-CTDx3-TEV-Kin28-HAx3

MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESRLERL
EQLFLLIFPREDLDMILKMDSLQDIKALLTGLFVQDNVNKDAVTDRLASVETDMPLTLRQHRIS
ATSSSEESSNKGQRQLTVSNYLFDDEDTPPNPKKEIELENLYFQGEFQYSPTSPSYSLTSPSYS
PTSPSLTTMAMKVNMEYTKEKKVGEGTYAVVYLGCQHSTGRKIAIKEIKTSEFKDGLDMSAIRE
VKYLQEMQHPNVIELIDIFMAYDNLNLVLEFLPTDLEVVIKDKSILFTPADIKAWMLMTLRGVY
HCHRNFILHRDLKPNNLLFSPDGQIKVADFGLARAIPAPHEILTSNVVTRWYRAPELLFGAKHY
TSAIDIWSVGVIFAELMLRIPYLPGQNDVDQMEVTFRALGTPTDRDWPEVSSFMTYNKLQIYPP
PSRDELRKRFIAASEYALDFMCGMLTMNPQKRWTAVQCLESDYFKELPPPSDPSSIKIRNVMAI
PGGGRIFYPYDVPDYAGYPYDVPDYAGSYPYDVPDYAAQCGRSS

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; ### from DNA Strider Thursday, February 5, 2004 12:10:50 PM ; DNA sequence pMK500 w/ TEV 8330 b.p. complete sequence
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GTCCACCAGCTAACATAAAATGTAAGCTCTCGGGGCTCTCTTGCCTTCCAACCCAGTCAGAAAT CGAGTTCCAATCCAAAAGTTCACCTGTCCCACCTGCTTCTGAATCAAACAAGGGAATAAACGAA TGAGGTTTCTGTGAAGCTGCACTGAGTAGTATGTTGCAGTCTTTTGGAAATACGAGTCTTTTAA TAACTGGCAAACCGAGGAACTCTTGGTATTCTTGCCACGACTCATCTCCATGCAGTTGGACGAT ATCAATGCCGTAATCATTGACCAGAGCCAAAACATCCTCCTTAGGTTGATTACGAAACACGCCA ACCAAGTATTTCGGAGTGCCTGAACTATTTTTATATGCTTTTACAAGACTTGAAATTTTCCTTG CAATAACCGGGTCAATTGTTCTCTTTCTATTGGGCACACATATAATACCCAGCAAGTCAGCATC $\tt GGAATCTAGTGCACATTCTGCGGCCTCTGTGCTCTGCAAGCCGCAAACTTTCACCAATGGACCA$ GAACTACCTGTGAAATTAATAACAGACATactccaagctgcctttgtgtgcttaatcacgtata atta attcttga agacga aagggcctcgtgatacgcctatttttataggttaatgtcatgataa ${\tt taatggtttcttagacgtcaggtggcacttttcggggaaatgtgcgcggaacccctatttgttt}$ atttttctaaatacattcaaatatgtatccgctcatgagacaataaccctgataaatgcttcaa ${\tt taatattgaaaaaggaagagtATGAGTATTCAACATTTCCGTGTCGCCCTTATTCCCTTTTTTG}$ $\tt CGGCATTTTGCCTCTGTTTTTGCTCACCCAGAAACGCTGGTGAAAGTAAAAGATGCTGAAGA$ ${\tt TCAGTTGGGTGCACGAGTGGGTTACATCGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAGT}$ $\tt TTTCGCCCCGAAGAACGTTTTCCAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTAT$ TATCCCGTATTGACGCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACTT ${\tt GGTTGAGTACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATGC}$ $\tt CGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACTCGCCTTGATCGTTGGGA$ ACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGACACCACGATGCCTGTAGCAATGGCA ACAACGTTGCGCAAACTATTAACTGGCGAACTACTTACTCTAGCTTCCCGGCAACAATTAATAG TATTGCTGATAAATCTGGAGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCA GATGGTAAGCCCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCAACTATGGATGAAC GAAATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATTGGtaactgtcagaccaagt ttactcatatatactttagattgatttaaaacttcatttttaatttaaaaggatctaggtgaag atcctttttgataatctcatgaccaaaatcccttaacgtgagttttcgttccactgagcgtcag $\verb|tttccgaaggtaactggcttcagcagagcgcagataccaaatactgtccttctagtgtagccgt|\\$ agttaggccaccacttcaagaactctgtagcaccgcctacatacctcgctctgctaatcctgtt accagtggctgctgccagtggcgataagtcgtgtcttaccgggttggactcaagacgatagtta ccggataaggcgcagcggtcgggctgaacggggggttcgtgcacacagcccagcttggagcgaa cgacctacaccgaactgagatacctacagcgtgagctatgagaaagcgccacgcttcccgaagg $\tt gagaaaggcggacaggtatccggtaagcggcagggtcggaacaggagagcgcacgagggagctt$ ccagggggaaacgcctggtatctttatagtcctgtcgggtttcgccacctctgacttgagcgtc $\tt gatttttgtgatgctcgtcaggggggggggggcctatggaaaaacgccagcaacgcggccttttt$ acggttcctggccttttgctggccttttgctcacatgttctttcctgcgttatcccctgattct gcagcgagtcagtgagcgaggaagcggaagagcgcccaatacgcaaaccgcctctccccgcgcg ttggccgattcattaatgcaggatccgggatcgaagaaatgatggtaaatgaaataggaaatca tgatgtatttggctttgcggcgccgaaaaaacgagtttacgcaattgcacaatcatgctgactc tgtggcggacccgcgctcttgccggcccggcgataacgctgggcgtgaggctgtgcccggcgga $\tt gttttttgcgcctgcattttccaaggtttaccctgcgctaaggggcgagattggagaagcaata$

434-2104: Gal4 DBD-TEV-CTDx3-Kin28-HAx3
434-874: Gal4 DBD (amino acids 1-147)
932-952: TEV cleavage sequence (ENLYFQG)
962-1024: CTD (three tandem repeats of YSTPSPS)
1040-1957: Kin28 (cDNA) with E54Q mutation
1991-2080: trimeric HA

Translation: Gal4 DBD-CTDx3-TEV-Kin28-HAx3

MKLLSSIEQACDICRLKKLKCSKEKPKCAKCLKNNWECRYSPKTKRSPLTRAHLTEVESRLERL
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; ### from DNA Strider Thursday, February 5, 2004 12:10:50 PM; DNA sequence pMK502 w/ TEV 8330 b.p. complete sequence

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AACAAGATGGTATAGAGCGCCAGAATTGTTGTTTGGAGCTAAACATTACACATCGGCTATTGAT ATCTGGTCAGTAGGCGTTATATTCGCGGAATTAATGCTAAGGATACCTTATTTACCAGGACAGA ATGATGTCGATCAAATGGAAGTAACGTTCAGGGCCTTAGGGACACCTACAGATAGAGATTGGCC CGAAGTTTCTTCCTTTATGACGTATAACAAGTTACAAATATATCCGCCCCCTTCAAGAGATGAA TTGAGGAAAAGGTTCATTGCTGCTAGCGAATACGCCTTAGATTTTATGTGTGGAATGCTAACGA TGAACCCACAAAAGAGGTGGACCGCTGTTCAGTGTTTAGAAAGTGATTATTTCAAAGAATTACC ACCACCAAGTGACCCGTCTTCAATAAAAATACGTAACgtcatggCAATTCCCGGtggcggccgc atcttttacccatacgatgttcctgactatgcgggctatccctatgacgtcccggactatgcag gatcccccgataCCGTCGACCTGCAGAGATCTAtgaatcgtagatactgaaaaaccccgcaagt t cact t caact g t g cat c g t g caccat c t caat t t c t t t a t a cat c g t t t t g c c t t c t tttatgtaactatactcctctaagtttcaatcttggccatgtaacctctgatctatagaattttt taaatgactagaattaatgcccatcttttttttggacctaaattcttcatgaaaatatattacg agggcttattcagaagctttggacttcttcgccagaggtttggtcaagtctccaatcaaggttg $\verb|tcggcttgtctaecttgccagaaatttacgaaaagatggaaaagggtcaaatcgttggtagata|\\$ $\verb|cgttgttgacacttctaaataagcgaatttcttatgatttatgatttttattattaaataagtt|\\$ ataaaaaaaaataagtgtatacaaattttaaagtgactcttaggttttaaaacgaaaattcttat $\verb|tcttgagtaactctttcctgtaggtcaggttgctttctcaggtatagcatgaggtcgctcttat|\\$ tgaccacacctctaccggcatgccgagcaaatgcctgcaaatcgctccccatttcacccaattg tagatatgctaactccagcaatgagttgatgaatctcggtgtgtattttatgtcctcagaggac ${\tt aacacctgttgtaatcgttcttccacacggatcctggcgtaatagcgaagaggcccgcaccgat}$ $\verb|cgcccttcccaacagttgcgcagcctgaatggcgaatggcgcctgatgcggtattttctcctta|\\$ cgcatctgtgcggtatttcacaccgcatatatcgctgggccattctcatgaagaatatcttgaa $\verb|tttattgtcatattactagttggtggaagtccatatatcggtgatcaatatagtggttgaca|\\$ tgctggctagtcaacattgagccttttgatcatgcaaatatattacggtattttacaatcaaat atcaaacttaactattgactttataacttatttaggtggtaacattcttataaaaaagaaaaaa attactgcaaaacagtactagcttttaacttgtatcctaggttatctatgctgtctcaccatag agaatattacctatttcagaatgtatgtccatgattcgccgggtaaatacatataatacacaaa $\verb|ttttaagaaaatttcttttgactaagtccatatcgactttgtaaaagttcactttagcatacat|\\$ atattacacgagccagaaattgtaacttttgcctaaaatcacaaattgcaaaatttaattgctt tttaaacataaatgaaataatttatttattgtttatgattaccgaaacataaaacctgctcaag aaaaagaaactgttttgtccttggaaaaaaagcactacctaggagcggccaaaatgccgaggct ttcatagcttaaactctttacagaaaataggcattatagatcagttcgagttttcttattcttc $\verb"cttccggttttatcgtcacagttttacagtaaataagtatcacctcttagagttcgatgataag"$ ctgtcaaacatgagaattaattccacatgttaaaatagtgaaggagcatgttcggcacacagtg gaccgaacgtggggtaagtgcactagggtccggttaaacggatctcgcattgatgaggcaacgc taattatcaacatatagattgttatctatctgcatgaacacgaaatctttacttgacgacttga $\verb|ggctgatggtgtttatgcaaagaaaccactgtgtttaatatgtgtcactgtttgatattactgt|\\$ $\verb|cagcgtagaagataatagtaaaagcggttaataagtgtatttgagataagtgtgataaagtttt|\\$ tacagcgaaaagacgataaatacaagaaaatgattacgaggatacggagagaggtatgtacatg tgtatttatatactaagctgccggcggttgtttgcaagaccgagaaaaggctagcaagaatcgg $\tt gtcattgtagcgtatgcgcctgtgaacattctcttcaacaagtttgattccattgcggtgaaat$

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